GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: January 4, 2003, 23:31:50 ; Search time 47 Seconds
(without alignments)
6531.563 Million cell updates/sec

Title: US-09-740-211-14_COPY_2040_3040
Perfect score: 1001
Sequence: 1 aatcagatcctcggtgcctg.....ctcagcccttataccgtgga 1001

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1001	100.0	4999	4	US-09-470-618-14	Sequence 14, Appl
2	1001	100.0	4999	4	US-09-364-862-14	Sequence 14, Appl
3	947.4	94.6	4629	2	US-08-484-891-7	Sequence 7, Appli
4	947.4	94.6	4670	3	US-08-717-294-41	Sequence 41, Appl
5	947.4	94.6	9354	1	US-08-683-839B-2	Sequence 2, Appli
6	919	91.8	11933	4	US-09-470-618-13	Sequence 13, Appl
7	919	91.8	11933	4	US-09-364-862-13	Sequence 13, Appl
8	723.2	72.2	4404	4	US-09-523-656-37	Sequence 37, Appl
9	688.4	68.8	4334	2	US-08-670-707A-38	Sequence 38, Appl
10	688.4	68.8	4334	4	US-09-037-601-38	Sequence 38, Appl
11	688.4	68.8	4334	4	US-09-315-179-38	Sequence 38, Appl
12	671	67.0	6999	1	US-08-276-594A-1	Sequence 1, Appli
13	671	67.0	7056	1	US-08-121-202-1	Sequence 1, Appli
14	671	67.0	8241	6	5171844-1	Patent No. 5171844
15	671	67.0	8967	1	US-08-366-851A-1	Sequence 1, Appli
16	671	67.0	9009	1	US-07-864-004B-3	Sequence 3, Appli
17	671	67.0	9009	1	US-08-251-937A-3	Sequence 3, Appli
18	671	67.0	9009	1	US-08-212-133A-1	Sequence 1, Appli
19	671	67.0	9009	1	US-08-474-503-1	Sequence 1, Appli
20	671	67.0	9009	2	US-08-670-707A-1	Sequence 1, Appli
21	671	67.0	9009	4	US-09-037-601-1	Sequence 1, Appli
22	671	67.0	9009	4	US-09-315-179-1	Sequence 1, Appli
23	671	67.0	9009	4	US-09-523-656-1	Sequence 1, Appli
24	671	67.0	9009	5	PCT-US93-03275-3	Sequence 3, Appli
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26	575	57.4	7032	4	US-09-324-867-1	Sequence 1, Appli
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28	566	56.5	5035	2	US-08-558-107-1	Sequence 1, Appli
29	566	56.5	5035	3	US-09-243-539-1	Sequence 1, Appli
30	560.6	56.0	4451	3	US-08-717-294-42	Sequence 42, Appl
31	554.4	55.4	6402	2	US-08-670-707A-36	Sequence 36, Appl
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33	554.4	55.4	6402	4	US-09-315-179-36	Sequence 36, Appl
34	554.4	55.4	6402	4	US-09-523-656-29	Sequence 29, Appl
35	551	55.0	1130	1	US-07-864-004B-1	Sequence 1, Appli
36	551	55.0	1130	1	US-08-251-937A-1	Sequence 1, Appli
37	551	55.0	1130	1	US-08-212-133A-5	Sequence 5, Appli
38	551	55.0	1130	1	US-08-474-503-3	Sequence 3, Appli
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41	551	55.0	1130	4	US-09-315-179-3	Sequence 3, Appli
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43	551	55.0	1130	5	PCT-US93-03275-1	Sequence 1, Appli
44	551	55.0	1130	5	PCT-US94-13200-3	Sequence 3, Appli
45	541.4	54.1	7493	1	US-08-212-133A-7	Sequence 7, Appli

ALIGNMENTS

RESULT 1
US-09-470-618-14
; Sequence 14, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/09/470,618
; CURRENT FILING DATE: 1999-12-22
; EARLIER APPLICATION NUMBER: 09/364,862
; EARLIER FILING DATE: 1999-07-30
; EARLIER APPLICATION NUMBER: 60/125,974
; EARLIER FILING DATE: 1999-03-24
; EARLIER APPLICATION NUMBER: 60/104,994
; EARLIER FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 4999
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-470-618-14

Query Match 100.0%; Score 1001; DB 4; Length 4999;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
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Db    2100 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2159

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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2340 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2399

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
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Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
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Db   2580 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2639

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
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Db   2760 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2819

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Db   2940 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 2999

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Db   3000 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3040
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RESULT 2
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; Sequence 14, Application US/09364862
; Patent No. 6221349
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: ADENO-ASSOCIATED VECTORS FOR EXPRESSION OF FACTOR VIII
;  TITLE OF INVENTION: BY TARGET
;  TITLE OF INVENTION: CELLS
;  FILE REFERENCE: AVIGEN-03743
;  CURRENT APPLICATION NUMBER: US/09/364,862
;  CURRENT FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 14
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-364-862-14

  Query Match             100.0%;  Score 1001;  DB 4;  Length 4999;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db   2040 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2099

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
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Db   2340 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2399

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
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Db   2460 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2519

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Db   2580 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2639

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Db   2640 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2699

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
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Db   2760 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2819

Qy    781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
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Qy    841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
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Db    3000 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3040


RESULT 3
US-08-484-891-7
; Sequence 7, Application US/08484891
; Patent No. 5935935
;  GENERAL INFORMATION:
;    APPLICANT:  Connelly, Sheila
;    APPLICANT:  Kaleko, Michael
;    APPLICANT:  Smith, Theodore
;    TITLE OF INVENTION:  Adenoviral Vectors for
;    TITLE OF INVENTION:  Treatment of Hemophilia
;    NUMBER OF SEQUENCES:  7
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Carella, Byrne, Bain, Gilfillan,
;      ADDRESSEE:  Cecchi, Stewart & Olstein
;      STREET:  6 Becker Farm Road
;      CITY:  Roseland
;      STATE:  New Jersey
;      COUNTRY:  USA
;      ZIP:  07068
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3.5 inch diskette
;      COMPUTER:  IBM PS/2
;      OPERATING SYSTEM:  MS-DOS
;      SOFTWARE:  WordPerfect 5.1
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/484,891
;      FILING DATE:  07-JUN-1995
;      CLASSIFICATION:  514
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/218,335
;      FILING DATE:  25-MAR-1994
;      APPLICATION NUMBER:  08/074,920
;      FILING DATE:  10-JUN-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Olstein, Elliot M.
;      REGISTRATION NUMBER:  24,025
;      REFERENCE/DOCKET NUMBER:  271010-273
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  201-994-1700
;      TELEFAX:  201-994-1744
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  4629 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA primer
;    FEATURE:
;      NAME/KEY:  Factor VIII cDNA with
;      NAME/KEY:  B domain deleted
US-08-484-891-7

  Query Match             94.6%;  Score 947.4;  DB 2;  Length 4629;
  Best Local Similarity   95.9%;  Pred. No. 1.4e-300;
  Matches 1000;  Conservative    0;  Mismatches    1;  Indels   42;  Gaps    1;

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Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
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Db    1745 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1804

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Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
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Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
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Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
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Qy     679 TGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 738
           |||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db    2345 TGAAACGCCATCAACGGGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 2404

Qy     739 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 798
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2405 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 2464

Qy     799 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 858
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2465 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 2524

Qy     859 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 918
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2525 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 2584

Qy     919 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 978
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2585 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 2644

Qy     979 TTACTCAGCCCTTATACCGTGGA 1001
           |||||||||||||||||||||||
Db    2645 TTACTCAGCCCTTATACCGTGGA 2667


RESULT 4
US-08-717-294-41
; Sequence 41, Application US/08717294
; Patent No. 6114148
;  GENERAL INFORMATION:
;    APPLICANT:  SEED, BRIAN
;    APPLICANT:  HAAS, JURGEN
;    TITLE OF INVENTION:  HIGH LEVEL EXPRESSION OF
;    TITLE OF INVENTION:  PROTEINS
;    NUMBER OF SEQUENCES:  110
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Clark & Elbing LLP
```

```
;      STREET:  176 Federal Street
;      CITY:  Boston
;      STATE:  MA
;      COUNTRY:  USA
;      ZIP:  02110
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette
;      COMPUTER:  IBM Compatible
;      OPERATING SYSTEM:  DOS
;      SOFTWARE:  FastSEQ for Windows Version 2.0
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/717,294
;      FILING DATE:  20-SEP-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Elbing, Karen L
;      REGISTRATION NUMBER:  35,238
;      REFERENCE/DOCKET NUMBER:  00786/345001
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  617-428-0200
;      TELEFAX:  617-428-7045
;      TELEX:
;  INFORMATION FOR SEQ ID NO:  41:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  4670 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
US-08-717-294-41

  Query Match             94.6%;  Score 947.4;  DB 3;  Length 4670;
  Best Local Similarity   95.9%;  Pred. No. 1.4e-300;
  Matches 1000;  Conservative    0;  Mismatches    1;  Indels   42;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1651 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1710

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1711 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1770

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1771 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1830

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1831 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 1890

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1891 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 1950

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1951 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2010

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2011 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2070

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2071 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2130

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2131 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2190

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2191 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2250

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2251 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2310

Qy     661 CCCAGAAT----------------------------------------CCACCAGTCT 678
          ||||||||                                        ||||||||||
Db    2311 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCCCACCAGTCT 2370

Qy     679 TGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 738
          |||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db    2371 TGAAACGCCATCAACGGGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 2430

Qy     739 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 798
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2431 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 2490

Qy     799 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 858
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2491 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 2550

Qy     859 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 918
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2551 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 2610

Qy     919 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 978
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2611 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 2670

Qy     979 TTACTCAGCCCTTATACCGTGGA 1001
          |||||||||||||||||||||||
Db    2671 TTACTCAGCCCTTATACCGTGGA 2693


RESULT 5
US-08-683-839B-2
; Sequence 2, Application US/08683839B
; Patent No. 5744326
;  GENERAL INFORMATION:
;    APPLICANT:  Ill, Charles . R. et al.
;    TITLE OF INVENTION:  Use of Viral Cis-Acting Post-Transcriptional
;    TITLE OF INVENTION:  Regulatory Sequences To Increase Expression of
;    TITLE OF INVENTION:  Intronless Genes Containing Near-Consensus Splice Sites
;    NUMBER OF SEQUENCES:  18
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  LAHIVE & COCKFIELD
;      STREET:  60 State Street, suite 510
;      CITY:  Boston
;      STATE:  Massachusetts
;      COUNTRY:  USA
;      ZIP:  02109-1875
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/683,839B
;      FILING DATE:  11-MARCH-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Remillard, Jane E.
;      REGISTRATION NUMBER:  38,872
;      REFERENCE/DOCKET NUMBER:  TTI-138
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)227-7400
```

```
;      TELEFAX:  (617)227-5941
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9354 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  2965..7378
US-08-683-839B-2

  Query Match             94.6%;  Score 947.4;  DB 1;  Length 9354;
  Best Local Similarity   95.9%;  Pred. No. 2.1e-300;
  Matches 1000;  Conservative    0;  Mismatches    1;  Indels   42;  Gaps    1;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4589 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 4648

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4649 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 4708

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4709 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 4768

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4769 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 4828

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4829 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 4888

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4889 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 4948

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4949 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 5008

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5009 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 5068

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5069 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 5128

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5129 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 5188

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5189 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 5248

Qy    661 CCCAGAAT----------------------------------------CCACCAGTCT 678
          ||||||||                                        ||||||||||
Db   5249 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCCCACCAGTCT 5308

Qy    679 TGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 738
          |||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||
Db   5309 TGAAACGCCATCAACGGGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 5368

Qy    739 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 798
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5369 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 5428

Qy    799 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 858
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5429 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 5488

Qy    859 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 918
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5489 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 5548

Qy    919 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 978
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5549 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 5608

Qy    979 TTACTCAGCCCTTATACCGTGGA 1001
          |||||||||||||||||||||||
Db   5609 TTACTCAGCCCTTATACCGTGGA 5631


RESULT 6
US-09-470-618-13
; Sequence 13, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION: by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/470,618
;  CURRENT FILING DATE: 1999-12-22
;  EARLIER APPLICATION NUMBER: 09/364,862
;  EARLIER FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-470-618-13

  Query Match             91.8%;  Score 919;  DB 4;  Length 11933;
  Best Local Similarity   96.4%;  Pred. No. 5.2e-291;
  Matches  965;  Conservative    0;  Mismatches    0;  Indels   36;  Gaps    1;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2051 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2110

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2111 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2170

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2171 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 2230

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2231 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2290

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2291 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2350

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2351 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2410
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Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2411 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2470

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2471 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2530

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2531 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2590

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2591 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2650

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2651 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTT-- 2708

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
           ||||||||||||||||||||||||||||||
Db    2709 ------------------------------CGAAATAACTCGTACTACTCTTCAGT 2734

Qy     721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2735 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2794

Qy     781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2795 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 2854

Qy     841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2855 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 2914

Qy     901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2915 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 2974

Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 1001
           |||||||||||||||||||||||||||||||||||||||||
Db    2975 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3015
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RESULT 7
US-09-364-862-13
; Sequence 13, Application US/09364862
; Patent No. 6221349
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: ADENO-ASSOCIATED VECTORS FOR EXPRESSION OF FACTOR VIII
; TITLE OF INVENTION: BY TARGET
; TITLE OF INVENTION: CELLS
; FILE REFERENCE: AVIGEN-03743
; CURRENT APPLICATION NUMBER: US/09/364,862
; CURRENT FILING DATE: 1999-07-30
; EARLIER APPLICATION NUMBER: 60/125,974
; EARLIER FILING DATE: 1999-03-24
; EARLIER APPLICATION NUMBER: 60/104,994
; EARLIER FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
; LENGTH: 11933
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic
US-09-364-862-13

Query Match 91.8%; Score 919; DB 4; Length 11933;

Best Local Similarity 96.4%; Pred. No. 5.2e-291;
Matches 965; Conservative 0; Mismatches 0; Indels 36; Gaps 1;

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Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2051 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2110

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2111 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2170

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2171 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 2230

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2231 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2290

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2291 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2350

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2351 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2410

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2411 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2470

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2471 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2530

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2531 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2590

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2591 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2650

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2651 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTT-- 2708

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
           ||||||||||||||||||||||||||||||
Db    2709 ------------------------------CGAAATAACTCGTACTACTCTTCAGT 2734

Qy     721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2735 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2794

Qy     781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2795 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 2854

Qy     841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2855 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 2914

Qy     901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2915 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 2974

Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 1001
           |||||||||||||||||||||||||||||||||||||||||
Db    2975 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3015
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RESULT 8

```
US-09-523-656-37
; Sequence 37, Application US/09523656
; Patent No. 6458563
; GENERAL INFORMATION:
;  APPLICANT: Lollar S., John
;  TITLE OF INVENTION: MODIFIED FACTOR VIII
;  FILE REFERENCE: 75-95I
;  CURRENT APPLICATION NUMBER: US/09/523,656
;  CURRENT FILING DATE: 2000-03-10
;  EARLIER APPLICATION NUMBER: 09/037,601
;  EARLIER FILING DATE: 1998-03-10
;  EARLIER APPLICATION NUMBER: 08/670,707
;  EARLIER FILING DATE: 1996-06-26
;  NUMBER OF SEQ ID NOS: 38
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 37
;   LENGTH: 4404
;   TYPE: DNA
;   ORGANISM: Porcine
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(4401)
US-09-523-656-37

  Query Match             72.2%;  Score 723.2;  DB 4;  Length 4404;
  Best Local Similarity   82.7%;  Pred. No. 5.5e-227;
  Matches  852;  Conservative    0;  Mismatches  148;  Indels   30;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           | || ||||||||||||||||||||| ||||| || | | ||||| | |||| |||||
Db    1625 AGTCCGATCCTCGGTGCCTGACCCGCTACTACTCGAGCTCCATTAATCTAGAGAAAGATC 1684

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | ||||| ||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db    1685 TGGCTTCGGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGACCAAAGAG 1744

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||||||||| |||||||||||||| || |||||||||||||||||||| |||||||| |
Db    1745 GAAACCAGATGATGTCAGACAAGAGAAACGTCATCCTGTTTTCTGTATTCGATGAGAATC 1804

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           |||||||||||||| |||||||||| || ||||| |||||||||| | |||| | ||||
Db    1805 AAAGCTGGTACCTCGCAGAGAATATTCAGCGCTTCCTCCCCAATCCGGATGGATTACAGC 1864

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
              |||||||||||||||||| || ||||||||||||||||||||||||||||||||||
Db    1865 CCCAGGATCCAGAGTTCCAAGCTTCTAACATCATGCACAGCATCAATGGCTATGTTTTTG 1924

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           |||| |||||| |||| |||||||||| |||||||||||||||||||||||||||||  |||
Db    1925 ATAGCTTGCAGCTGTCGGTTTGTTTGCACGAGGTGGCATACTGGTACATTCTAAGTGTTG 1984

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           |||||||||| |||||||| || |||||||||||||| || ||||||||||||||||||
Db    1985 GAGCACAGACGGACTTCCTCTCCGTCTTCTTCTCTGGCTACACCTTCAAACACAAAATGG 2044

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           |||||||||||||||||||||| ||||| |||||||||||||| |||||||||| ||||
Db    2045 TCTATGAAGACACACTCACCCTGTTCCCCTTCTCAGGAGAAACGGTCTTCATGTCAATGG 2104

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           |||||||||||||| ||| | || |||||||||||||||||| |||||||||| ||||
Db    2105 AAAACCCAGGTCTCTGGGTCCTTGGGTGCCACAACTCAGACTTGCGGAACAGAGGGATGA 2164

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           | |||||||||||||| | |||||||||| | ||| |||||||||||| ||  ||| ||
Db    2165 CAGCCTTACTGAAGGTGTATAGTTGTGACAGGGACATTGGTGATTATTATGACAACACTT 2224

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           |||||||||| |||  | |||||||||  ||| |||| ||||||||| || |||||
Db    2225 ATGAAGATATTCCAGGCTTCTTGCTGAGTGGAAAGAATGTCATTGAACCTAGGAGCTTTG 2284

Qy     661 CCCAGAAT----------------------------CCACCAGTCTTGAAACGCCATC 690
           |||||||                            || || ||| ||  ||| ||||
Db    2285 CCCAGAATTCAAGACCCCCTAGTGCGAGCGCTCCAAAGCCTCCGGTCCTGCGACGGCATC 2344

Qy     691 AACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTGACTATGATGATA 750
           |  | || ||||  | | ||||| ||||| | ||  ||||  |||| ||||||||||||
Db    2345 AGAGGGACATAAGCCTTCCTACTTTTCAGCCGGAGGAAGACAAAATGGACTATGATGATA 2404

Qy     751 CCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGGATGAAAATCAGA 810
           | | |||  ||||| ||||   |||||||||||||||| | ||||||||||||||||
Db    2405 TCTTCTCAACTGAAACGAAGGGAGAAGATTTTGACATTTACGGTGAGGATGAAAATCAGG 2464

Qy     811 GCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAGTGGAGAGGCTCT 870
           ||| |||||||||||| |||| ||| |||||||||||| |||||||| ||||||  |||||
Db    2465 ACCCTCGCAGCTTTCAGAAGAGAACCCGACACTATTTCATTGCTGCGGTGGAGCAGCTCT 2524

Qy     871 GGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTCAGAGTGGCAGTG 930
           ||||||| ||||||||    ||||| |  |  ||||||||||||||||||  ||    |
Db    2525 GGGATTACGGGATGAGCGAATCCCCCCGGGCGCTAAGAAACAGGGCTCAGAACGGAGAGG 2584

Qy     931 TCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCTTTACTCAGCCCT 990
           | |||| |||||||||| || || |||| ||||||| |||| |||||||| || ||||| |
Db    2585 TGCCTCGGTTCAAGAAGGTGGTCTTCCGGGAATTTGCTGACGGCTCCTTCACGCAGCCGT 2644

Qy     991 TATACCGTGG 1000
             ||||| ||
Db    2645 CGTACCGCGG 2654


RESULT 9
US-08-670-707A-38
; Sequence 38, Application US/08670707A
; Patent No. 5859204
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/670,707A
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Greenlee, Lorance L.
;      REGISTRATION NUMBER:  27,894
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  38:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  4334 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
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;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  cDNA to mRNA
;    HYPOTHETICAL:  NO
;    ORIGINAL SOURCE:
;      INDIVIDUAL ISOLATE:  Factor VIII lacking B domain
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  3..4334
US-08-670-707A-38

  Query Match             68.8%;  Score 688.4;  DB 2;  Length 4334;
  Best Local Similarity   82.2%;  Pred. No. 1.4e-215;
  Matches  822;  Conservative    0;  Mismatches  136;  Indels   42;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           | || ||||||||||||||||||||||| ||||| || | | ||||| | |||| |||||
Db    1627 AGTCCGATCCTCGGTGCCTGACCCGCTACTACTCGAGCTCCATTAATCTAGAGAAAGATC 1686

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | ||||| |||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db    1687 TGGCTTCGGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGACCAAAGAG 1746

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||||||||| |||||||||||||| || ||||||||||||||||||||| |||||||| |
Db    1747 GAAACCAGATGATGTCAGACAAGAGAAACGTCATCCTGTTTTCTGTATTCGATGAGAATC 1806

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           |||||||||||| |||||||||| || ||||| |||||||||| | |||| | ||||
Db    1807 AAAGCTGGTACCTCGCAGAGAATATTCAGCGCTTCCTCCCCAATCCGGATGGATTACAGC 1866

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||||||||||||||||||| || ||||||||||||||||||||||||||||||||||||
Db    1867 CCCAGGATCCAGAGTTCCAAGCTTCTAACATCATGCACAGCATCAATGGCTATGTTTTTG 1926

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           |||| |||||| |||| |||||||||| |||||||||||||||||||||||||  |||
Db    1927 ATAGCTTGCAGCTGTCGGTTTGTTTGCACGAGGTGGCATACTGGTACATTCTAAGTGTTG 1986

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           |||||||||| |||||||| || |||||||||||||| || ||||||||||||||||||
Db    1987 GAGCACAGACGGACTTCCTCTCCGTCTTCTTCTCTGGCTACACCTTCAAACACAAAATGG 2046

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           |||||||||||||||||||| ||||| |||||||||||||| |||||||||| ||||
Db    2047 TCTATGAAGACACACTCACCCTGTTCCCCTTCTCAGGAGAAACGGTCTTCATGTCAATGG 2106

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           |||||||||||| ||| | || |||||||||||||||||||| |||||||||| ||||
Db    2107 AAAACCCAGGTCTCTGGGTCCTAGGGTGCCACAACTCAGACTTGCGGAACAGAGGGATGA 2166

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           | |||||||||||||| | |||||||||| | ||| |||||||||||  ||  ||| ||
Db    2167 CAGCCTTACTGAAGGTGTATAGTTGTGACAGGGACATTGGTGATTATTATGACAACACTT 2226

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           |||||||||| |||  | |||||||||  ||| |||| ||||||||| |||
Db    2227 ATGAAGATATTCCAGGCTTCTTGCTGAGTGGAAAGAATGTCATTGAACCCAGA------- 2279

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
                                         || ||||  | | ||||| |||||
Db    2280 ----------------------------------GACATAAGCCTTCCTACTTTTCAGC 2304

Qy     721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
           | ||  ||||  |||| |||||||||||| | | |||  ||||| ||||  |||||||
Db    2305 CGGAGGAAGACAAAATGGACTATGATGATATCTTCTCAACTGAAACGAAGGGAGAAGATT 2364

Qy     781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
           |||||||||| | |||||||||||||||  ||| |||||||||| |||| ||| ||||
Db    2365 TTGACATTTACGGTGAGGATGAAAATCAGGACCCTCGCAGCTTTCAGAAGAGAACCCGAC 2424

Qy     841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
           ||||||| |||||||| |||||| |||||||||||| ||||||||    ||||| |  |
Db    2425 ACTATTTCATTGCTGCGGTGGAGCAGCTCTGGGATTACGGGATGAGCGAATCCCCCCGGG 2484

Qy     901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
           ||||||||||||||||||||  ||    || |||| |||||||| || || |||| ||
Db    2485 CGCTAAGAAACAGGGCTCAGAACGGAGAGGTGCCTCGGTTCAAGAAGGTGGTCTTCCGGG 2544

Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGG 1000
           ||||| |||| |||||||| || ||||| |  ||||| ||
Db    2545 AATTTGCTGACGGCTCCTTCACGCAGCCGTCGTACCGCGG 2584


RESULT 10
US-09-037-601-38
; Sequence 38, Application US/09037601
; Patent No. 6180371
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/037,601
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Ferber, Donna M.
;      REGISTRATION NUMBER:  33,878
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  38:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  4334 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  cDNA to mRNA
;    HYPOTHETICAL:  NO
;    ORIGINAL SOURCE:
;      INDIVIDUAL ISOLATE:  Factor VIII lacking B domain
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  3..4334
US-09-037-601-38

  Query Match             68.8%;  Score 688.4;  DB 4;  Length 4334;
  Best Local Similarity   82.2%;  Pred. No. 1.4e-215;
  Matches  822;  Conservative    0;  Mismatches  136;  Indels   42;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           | || ||||||||||||||||||||||| ||||| || | | ||||| | |||| |||||
Db    1627 AGTCCGATCCTCGGTGCCTGACCCGCTACTACTCGAGCTCCATTAATCTAGAGAAAGATC 1686
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Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db    1687 TGGCTTCGGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGACCAAAGAG 1746

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||||||||| |||||||||||||| || ||||||||||||||||||||| |||||||| |
Db    1747 GAAACCAGATGATGTCAGACAAGAGAAACGTCATCCTGTTTTCTGTATTCGATGAGAATC 1806

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           |||||||||||||| |||||||||| || ||||| |||||||||| | |||| | ||||
Db    1807 AAAGCTGGTACCTCGCAGAGAATATTCAGCGCTTCCTCCCCAATCCGGATGGATTACAGC 1866

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||||||||||||||||||| || ||||||||||||||||||||||||||||||||||||||
Db    1867 CCCAGGATCCAGAGTTCCAAGCTTCTAACATCATGCACAGCATCAATGGCTATGTTTTTG 1926

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           |||| |||||| |||| |||||||||| |||||||||||||||||||||||||||  |||
Db    1927 ATAGCTTGCAGCTGTCGGTTTGTTTGCACGAGGTGGCATACTGGTACATTCTAAGTGTTG 1986

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           |||||||||| |||||||| || |||||||||||||| || ||||||||||||||||||||
Db    1987 GAGCACAGACGGACTTCCTCTCCGTCTTCTTCTCTGGCTACACCTTCAAACACAAAATGG 2046

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           |||||||||||||||||||| ||||| |||||||||||||| |||||||||| ||||
Db    2047 TCTATGAAGACACACTCACCCTGTTCCCCTTCTCAGGAGAAACGGTCTTCATGTCAATGG 2106

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           |||||||||||||| ||| | || |||||||||||||||||||| |||||||||| ||||
Db    2107 AAAACCCAGGTCTCTGGGTCCTAGGGTGCCACAACTCAGACTTGCGGAACAGAGGGATGA 2166

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           | |||||||||||||| | |||||||||| | ||| |||||||||||  ||  ||| ||
Db    2167 CAGCCTTACTGAAGGTGTATAGTTGTGACAGGGACATTGGTGATTATTATGACAACACTT 2226

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           |||||||||| |||  | |||||||||  ||| |||| |||||||||  |||
Db    2227 ATGAAGATATTCCAGGCTTCTTGCTGAGTGGAAAGAATGTCATTGAACCCAGA------- 2279

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
                                          || ||||  | | ||||| |||||
Db    2280 ----------------------------------GACATAAGCCTTCCTACTTTTCAGC 2304

Qy     721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
           | ||  ||||  |||| |||||||||||||| | | |||  ||||| |||   |||||||
Db    2305 CGGAGGAAGACAAAATGGACTATGATGATATCTTCTCAACTGAAACGAAGGGAGAAGATT 2364

Qy     781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
           |||||||||| | |||||||||||||||  ||| |||||||||| |||| ||| ||||
Db    2365 TTGACATTTACGGTGAGGATGAAAATCAGGACCCTCGCAGCTTTCAGAAGAGAACCCGAC 2424

Qy     841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
           ||||||| |||||||| |||||| |||||||||||| ||||||||    ||||| | |
Db    2425 ACTATTTCATTGCTGCGGTGGAGCAGCTCTGGGATTACGGGATGAGCGAATCCCCCCGGG 2484

Qy     901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
             ||||||||||||||||||  ||    || |||| ||||||||| || || |||| ||
Db    2485 CGCTAAGAAACAGGGCTCAGAACGGAGAGGTGCCTCGGTTCAAGAAGGTGGTCTTCCGGG 2544

Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGG 1000
           ||||| |||| |||||||| || ||||| |  ||||| ||
Db    2545 AATTTGCTGACGGCTCCTTCACGCAGCCGTCGTACCGCGG 2584


RESULT 11
US-09-315-179-38
; Sequence 38, Application US/09315179
; Patent No. 6376463
; GENERAL INFORMATION:
;  APPLICANT: Lollar, John S
;  TITLE OF INVENTION: Modified Factor VIII
;  FILE REFERENCE: 75-95H
;  CURRENT APPLICATION NUMBER: US/09/315,179
;  CURRENT FILING DATE: 1999-05-20
;  EARLIER APPLICATION NUMBER: U.S. 09/037,601
;  EARLIER FILING DATE: 1998-03-10
;  EARLIER APPLICATION NUMBER: U.S. 08/670,707
;  EARLIER FILING DATE: 1996-06-26
;  EARLIER APPLICATION NUMBER: PCT/US97/11155
;  EARLIER FILING DATE: 1997-06-26
;  EARLIER APPLICATION NUMBER: PCT/US94/13200
;  EARLIER FILING DATE: 1994-11-15
;  EARLIER APPLICATION NUMBER: U.S. 08/212,133
;  EARLIER FILING DATE: 1994-03-11
;  EARLIER APPLICATION NUMBER: U.S. 07/864,004
;  EARLIER FILING DATE: 1992-04-07
;  NUMBER OF SEQ ID NOS: 40
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 38
;   LENGTH: 4334
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:factor VIII
;   OTHER INFORMATION: lacking the B domain
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (3)..(4331)
US-09-315-179-38

  Query Match             68.8%;  Score 688.4;  DB 4;  Length 4334;
  Best Local Similarity   82.2%;  Pred. No. 1.4e-215;
  Matches  822;  Conservative    0;  Mismatches  136;  Indels   42;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           | || |||||||||||||||||||||| ||||| || | | ||||| | |||| |||||
Db    1627 AGTCCGATCCTCGGTGCCTGACCCGCTACTACTCGAGCTCCATTAATCTAGAGAAAGATC 1686

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | ||||| |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db    1687 TGGCTTCGGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGACCAAAGAG 1746

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||||||||| |||||||||||||| || ||||||||||||||||||||| |||||||| |
Db    1747 GAAACCAGATGATGTCAGACAAGAGAAACGTCATCCTGTTTTCTGTATTCGATGAGAATC 1806

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           |||||||||||||| |||||||||| || ||||| |||||||||| | |||| | ||||
Db    1807 AAAGCTGGTACCTCGCAGAGAATATTCAGCGCTTCCTCCCCAATCCGGATGGATTACAGC 1866

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||||||||||||||||||| || ||||||||||||||||||||||||||||||||||||||
Db    1867 CCCAGGATCCAGAGTTCCAAGCTTCTAACATCATGCACAGCATCAATGGCTATGTTTTTG 1926

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           |||| |||||| |||| |||||||||| |||||||||||||||||||||||||||  |||
Db    1927 ATAGCTTGCAGCTGTCGGTTTGTTTGCACGAGGTGGCATACTGGTACATTCTAAGTGTTG 1986

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           |||||||||| |||||||| || |||||||||||||| || ||||||||||||||||||||
Db    1987 GAGCACAGACGGACTTCCTCTCCGTCTTCTTCTCTGGCTACACCTTCAAACACAAAATGG 2046

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           |||||||||||||||||||| ||||| |||||||||||||| |||||||||| ||||
Db    2047 TCTATGAAGACACACTCACCCTGTTCCCCTTCTCAGGAGAAACGGTCTTCATGTCAATGG 2106

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           |||||||||||||| ||| | || |||||||||||||||||||| |||||||||| ||||
Db    2107 AAAACCCAGGTCTCTGGGTCCTAGGGTGCCACAACTCAGACTTGCGGAACAGAGGGATGA 2166

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           | |||||||||||||| | |||||||||| | ||| |||||||||||  ||  ||| ||
Db    2167 CAGCCTTACTGAAGGTGTATAGTTGTGACAGGGACATTGGTGATTATTATGACAACACTT 2226

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
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Db   2227 ATGAAGATATTCCAGGCTTCTTGCTGAGTGGAAAGAATGTCATTGAACCCAGA------- 2279

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720

Db   2280 ----------------------------------GACATAAGCCTTCCTACTTTTCAGC 2304

Qy    721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780

Db   2305 CGGAGGAAGACAAAATGGACTATGATGATATCTTCTCAACTGAAACGAAGGGAGAAGATT 2364

Qy    781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840

Db   2365 TTGACATTTACGGTGAGGATGAAAATCAGGACCCTCGCAGCTTTCAGAAGAGAACCCGAC 2424

Qy    841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900

Db   2425 ACTATTTCATTGCTGCGGTGGAGCAGCTCTGGGATTACGGGATGAGCGAATCCCCCCGGG 2484

Qy    901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960

Db   2485 CGCTAAGAAACAGGGCTCAGAACGGAGAGGTGCCTCGGTTCAAGAAGGTGGTCTTCCGGG 2544

Qy    961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGG 1000

Db   2545 AATTTGCTGACGGCTCCTTCACGCAGCCGTCGTACCGCGG 2584
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RESULT 12
US-08-276-594A-1
; Sequence 1, Application US/08276594A
; Patent No. 5693499
;  GENERAL INFORMATION:
;    APPLICANT:  YONEMURA, Hiroshi
;    APPLICANT:  TAJIMA, Yoshitaka
;    APPLICANT:  SUGAWARA, Keishin
;    APPLICANT:  MASUDA, Kenichi
;    TITLE OF INVENTION:  PROCESS FOR PREPARING HUMAN COAGULATION
;    TITLE OF INVENTION:  FACTOR VIII PROTEIN COMPLEX
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/276,594A
;      FILING DATE:  18-JUL-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/950,191
;      FILING DATE:  24-SEP-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 243262/1991
;      FILING DATE:  24-SEP-1991
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  WEGNER, Harold C.
;      REGISTRATION NUMBER:  25,258
;      REFERENCE/DOCKET NUMBER:  74129/195/AOPA
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  6999 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..6996
US-08-276-594A-1

  Query Match             67.0%;  Score 671;  DB 1;  Length 6999;
  Best Local Similarity   94.6%;  Pred. No. 1e-209;
  Matches  695;  Conservative    0;  Mismatches   40;  Indels    0;  Gaps    0;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60

Db    1568 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1627

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120

Db    1628 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1687

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180

Db    1688 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1747

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240

Db    1748 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 1807

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300

Db    1808 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 1867

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360

Db    1868 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 1927

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420

Db    1928 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 1987

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480

Db    1988 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2047

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540

Db    2048 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2107

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600

Db    2108 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2167

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660

Db    2168 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2227

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720

Db    2228 CCCAGAATTCAAGACACCGTAGCACTAGGCAAAAGCAATTTAATGCCACCACAATTCCAG 2287

Qy     721 CAGATCAAGAGGAAA 735

Db    2288 AAAATGACATAGAGA 2302
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RESULT 13
US-08-121-202-1
; Sequence 1, Application US/08121202
; Patent No. 5563045
;  GENERAL INFORMATION:
;    APPLICANT:  Pittman, Debra
;    APPLICANT:  Rehemtulla, Alnawaz
;    APPLICANT:  Wozney, John M.
```

```
;    APPLICANT:  Kaufman, Randal J.
;    TITLE OF INVENTION:  CHIMERIC PROCOAGULANT PROTEINS
;    NUMBER OF SEQUENCES:  27
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Legal Affairs, Genetics Institute, Inc.
;      STREET:  87 CambridgePark Drive
;      CITY:  Cambridge
;      STATE:  MA
;      COUNTRY:  USA
;      ZIP:  02140
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/121,202
;      FILING DATE:  14-SEP-1993
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Meinert, M. C.
;      REGISTRATION NUMBER:  31,544
;      REFERENCE/DOCKET NUMBER:  GI 5195A
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)876-1210 X8574
;      TELEFAX:  (617) 876-5851
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  7056 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..7053
US-08-121-202-1

  Query Match             67.0%;  Score 671;  DB 1;  Length 7056;
  Best Local Similarity   94.6%;  Pred. No. 1e-209;
  Matches  695;  Conservative    0;  Mismatches   40;  Indels    0;  Gaps    0;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1625 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1684

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1685 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1744

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1745 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1804

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1805 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 1864

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1865 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 1924

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1925 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 1984

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1985 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2044

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2045 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2104

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2105 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2164

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2165 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2224

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2225 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2284

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
          |||||||| ||  |  |   |        |||    || | | |   || ||  |||
Db   2285 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCACAATTCCAG 2344

Qy    721 CAGATCAAGAGGAAA 735
            | || |    || |
Db   2345 AAAATGACATAGAGA 2359


RESULT 14
5171844-1
;Patent No. 5171844
;    APPLICANT: VAN OOYEN, ALBERT J.J.;PANNEKOEK, HANS;VERBEET,
;MARTINUS P.;VAN LEEN, ROBERT W.
;    TITLE OF INVENTION: PROTEINS WITH FACTOR VIII ACTIVITY
;PROCESS FOR THEIR PREPARATION USING GENETICALLY-ENGINEERED CELLS
;AND PHARMACEUTICAL COMPOSITIONS CONTAINING THEM
;    NUMBER OF SEQUENCES: 12
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/205,226
;      FILING DATE: 10-JUN-1988
;SEQ ID NO:1:
;      LENGTH: 8241
5171844-1

  Query Match             67.0%;  Score 671;  DB 6;  Length 8241;
  Best Local Similarity   94.6%;  Pred. No. 1.1e-209;
  Matches  695;  Conservative    0;  Mismatches   40;  Indels    0;  Gaps    0;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1631 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1690

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1691 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1750

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1751 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1810

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1811 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 1870

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1871 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 1930

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1931 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 1990

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1991 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2050

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2051 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2110

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2111 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2170

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2171 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2230

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2231 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2290

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
          |||||||| ||  |  |   |        |||    || | | |   || ||  |||
Db   2291 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCACAATTCCAG 2350

Qy    721 CAGATCAAGAGGAAA 735
          | || |    || |
Db   2351 AAAATGACATAGAGA 2365


RESULT 15
US-08-366-851A-1
; Sequence 1, Application US/08366851A
; Patent No. 5681746
;  GENERAL INFORMATION:
;    APPLICANT:  Bodner, Mordechai
;    APPLICANT:  De Polo, Nicolas J.
;    APPLICANT:  Hsu, David Chi-Tang
;    APPLICANT:  Chang, Steven
;    TITLE OF INVENTION:  Retroviral Delivery of Full Length Factor VIII
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Viagene, Inc.
;      STREET:  11055 Roselle Street
;      CITY:  San Diego
;      STATE:  California
;      COUNTRY:  U.S.A.
;      ZIP:  92121
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/366,851A
;      FILING DATE:
;      CLASSIFICATION:  514
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Chambers, Daniel M.
;      REGISTRATION NUMBER:  34,561
;      REFERENCE/DOCKET NUMBER:  930049.438
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (619) 452-1288
;      TELEFAX:  (619) 452-2616
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  8967 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  110..7165
US-08-366-851A-1

  Query Match             67.0%;  Score 671;  DB 1;  Length 8967;
  Best Local Similarity   94.6%;  Pred. No. 1.2e-209;
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Job time : 131 secs
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GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

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(without alignments)
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 381593 seqs, 216252194 residues

Total number of hits satisfying chosen parameters: 763186

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	2	1001	100.0	4999	10	US-09-740-211-14	Sequence 14, Appl
	3	947.4	94.6	4629	10	US-09-150-811-7	GENERAL INFORMA
	4	947.4	94.6	7944	12	US-10-095-718-1	Sequence 1, Appli
	5	919	91.8	11933	9	US-10-007-968-13	Sequence 13, Appl
	6	919	91.8	11933	10	US-09-740-211-13	Sequence 13, Appl
	7	696.6	69.6	7914	12	US-10-095-718-3	Sequence 3, Appli
	8	671	67.0	9009	9	US-09-957-641-1	Sequence 1, Appli
	9	153.6	15.3	6909	10	US-09-880-107-2275	Sequence 2275, Ap
	10	124.4	12.4	3700	10	US-09-917-800A-1539	Sequence 1539, Ap
	11	117.4	11.7	3321	9	US-09-970-966-175	Sequence 175, App
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ALIGNMENTS

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RESULT 1
US-10-007-968-14
; Sequence 14, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-14

  Query Match           100.0%;  Score 1001;  DB 9;  Length 4999;
  Best Local Similarity  100.0%;  Pred. No. 1.3e-285;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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RESULT 2
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; Sequence 14, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/09/740,211
; CURRENT FILING DATE: 2000-12-18
; PRIOR APPLICATION NUMBER: 09/470,618
; PRIOR FILING DATE: 1999-12-22
; PRIOR APPLICATION NUMBER: 60/125,974
; PRIOR FILING DATE: 1999-03-24
; PRIOR APPLICATION NUMBER: 60/104,994
; PRIOR FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 4999
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-740-211-14

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Db   2580 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2639

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Db   2640 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2699

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
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RESULT 3
US-09-150-811-7
;  GENERAL INFORMATION:
;       APPLICANT: Connelly, Sheila
;                  Kaleko, Michael
;                  Smith, Theodore
;       TITLE OF INVENTION: Adenoviral Vectors for     Treatment of Hemophilia
;       NUMBER OF SEQUENCES: 7
;       CORRESPONDENCE ADDRESS:
;            ADDRESSEE: Carella, Byrne, Bain, Gilfillan,     Cecchi, Stewart & Olstein
;            STREET: 6 Becker Farm Road
;            CITY: Roseland
;            STATE: New Jersey
;            COUNTRY: USA
;            ZIP: 07068
;       COMPUTER READABLE FORM:
;            MEDIUM TYPE: 3.5 inch diskette
;            COMPUTER: IBM PS/2
;            OPERATING SYSTEM: MS-DOS
;            SOFTWARE: WordPerfect 5.1
;       CURRENT APPLICATION DATA:
;            APPLICATION NUMBER: US/09/150,811
;            FILING DATE: 10-Sep-1998
;            CLASSIFICATION: <Unknown>
;       PRIOR APPLICATION DATA:
;            APPLICATION NUMBER: 08/484,891
;            FILING DATE: 07-JUN-1995
;            APPLICATION NUMBER: 08/218,335
;            FILING DATE: 25-MAR-1994
;            APPLICATION NUMBER: 08/074,920
;            FILING DATE: 10-JUN-1993
;       ATTORNEY/AGENT INFORMATION:
;            NAME: Olstein, Elliot M.
;            REGISTRATION NUMBER: 24,025
;            REFERENCE/DOCKET NUMBER: 271010-440
;       TELECOMMUNICATION INFORMATION:
;            TELEPHONE: 973-994-1700
;            TELEFAX: 973-994-1744
;       SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-150-811-7

  Query Match             94.6%;  Score 947.4;  DB 10;  Length 4629;
  Best Local Similarity   95.9%;  Pred. No. 8.5e-270;
  Matches 1000;  Conservative    0;  Mismatches    1;  Indels   42;  Gaps    1;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1625 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1684

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1685 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1744

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1745 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1804

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1805 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 1864

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1865 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 1924

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1925 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 1984

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1985 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2044

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2045 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2104

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2105 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2164

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2165 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2224

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2225 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2284

Qy    661 CCCAGAAT----------------------------------------------CCACCAGTCT 678
          ||||||||                                              ||||||||||
Db   2285 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCCCACCAGTCT 2344

Qy    679 TGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 738
          |||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db   2345 TGAAACGCCATCAACGGGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 2404

Qy    739 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 798
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2405 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 2464

Qy    799 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 858
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2465 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 2524

Qy    859 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 918
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2525 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 2584

Qy    919 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 978
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2585 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 2644

Qy    979 TTACTCAGCCCTTATACCGTGGA 1001
          |||||||||||||||||||||||
Db   2645 TTACTCAGCCCTTATACCGTGGA 2667


RESULT 4
US-10-095-718-1
; Sequence 1, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
```

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;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
;  PRIOR FILING DATE: 2001-08-22
;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
;   LENGTH: 7944
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Plasmid pDLZ6 encoding Homo sapiens BDD FVIII
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (420)...(4835)
US-10-095-718-1

  Query Match             94.6%;  Score 947.4;  DB 12;  Length 7944;
  Best Local Similarity   95.9%;  Pred. No. 1.1e-269;
  Matches 1000;  Conservative    0;  Mismatches    1;  Indels   42;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2044 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2103

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2104 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2163

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2164 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 2223

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2224 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2283

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2284 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2343

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2344 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2403

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2404 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2463

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
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Db    2464 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2523

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
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Db    2524 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2583

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
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Db    2584 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2643

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2644 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2703

Qy     661 CCCAGAAT----------------------------------------CCACCAGTCT 678
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Db    2704 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCCCACCAGTCT 2763

Qy     679 TGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 738
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Db    2764 TGAAACGCCATCAACGGGAAATAACTCGTACTACTCTTCAGTCAGATCAAGAGGAAATTG 2823

Qy     739 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 798
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2824 ACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATTTTGACATTTATGATGAGG 2883

Qy     799 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 858
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Db    2884 ATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGACACTATTTTATTGCTGCAG 2943

Qy     859 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 918
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2944 TGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATGTTCTAAGAAACAGGGCTC 3003

Qy     919 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 978
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Db    3004 AGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGGAATTTACTGATGGCTCCT 3063

Qy     979 TTACTCAGCCCTTATACCGTGGA 1001
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Db    3064 TTACTCAGCCCTTATACCGTGGA 3086


RESULT 5
US-10-007-968-13
; Sequence 13, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-13

  Query Match             91.8%;  Score 919;  DB 9;  Length 11933;
  Best Local Similarity   96.4%;  Pred. No. 3.4e-261;
  Matches  965;  Conservative    0;  Mismatches    0;  Indels   36;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
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Db    2051 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2110

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
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Db    2111 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2170

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2171 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 2230

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2231 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2290

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
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Db   2291 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2350

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
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Db   2351 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2410

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2411 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2470

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2471 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2530

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2531 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2590

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2591 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2650

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2651 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTT-- 2708

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
                                          ||||||||||||||||||||||||||
Db   2709 ----------------------------------CGAAATAACTCGTACTACTCTTCAGT 2734

Qy    721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2735 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2794

Qy    781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
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Db   2795 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 2854

Qy    841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2855 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 2914

Qy    901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2915 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 2974

Qy    961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   2975 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3015
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RESULT 6
US-09-740-211-13
; Sequence 13, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-13

  Query Match             91.8%;  Score 919;  DB 10;  Length 11933;
  Best Local Similarity   96.4%;  Pred. No. 3.4e-261;
  Matches  965;  Conservative    0;  Mismatches    0;  Indels   36;  Gaps    1;

Qy      1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2051 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 2110

Qy     61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2111 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2170

Qy    121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2171 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 2230

Qy    181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2231 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2290

Qy    241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2291 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2350

Qy    301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2351 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2410

Qy    361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2411 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2470

Qy    421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2471 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2530

Qy    481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2531 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2590

Qy    541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2591 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2650

Qy    601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2651 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTT-- 2708

Qy    661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
                                          ||||||||||||||||||||||||||
Db   2709 ----------------------------------CGAAATAACTCGTACTACTCTTCAGT 2734

Qy    721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2735 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 2794

Qy    781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2795 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 2854

Qy    841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2855 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 2914

Qy    901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2915 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 2974
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Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 1001
           |||||||||||||||||||||||||||||||||||||||||||
Db    2975 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 3015
```

RESULT 7
US-10-095-718-3
; Sequence 3, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
; APPLICANT: Walsh, Christopher
; APPLICANT: Chao, Hengjun
; APPLICANT: Burstein, Haim
; APPLICANT: Lynch, Carmel
; APPLICANT: Stepan, Tony
; APPLICANT: Munson, Keith
; TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
; TITLE OF INVENTION: Methods of Using the Same
; FILE REFERENCE: 35052/204375
; CURRENT APPLICATION NUMBER: US/10/095,718
; CURRENT FILING DATE: 2002-03-12
; PRIOR APPLICATION NUMBER: 09/689,430
; PRIOR FILING DATE: 2001-08-22
; PRIOR APPLICATION NUMBER: 60/158,780
; PRIOR FILING DATE: 1999-10-12
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3
; LENGTH: 7914
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: rAAV vector with canine B-domain deleted factor
; OTHER INFORMATION: VIII
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (435)...(4730)
US-10-095-718-3

Query Match 69.6%; Score 696.6; DB 12; Length 7914;
Best Local Similarity 83.1%; Pred. No. 1.5e-195;
Matches 832; Conservative 0; Mismatches 109; Indels 60; Gaps 1;

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Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           |||||||||||||||||||||||| |||||||| || ||| ||||| ||||||||||||
Db    2041 AATCAGATCCTCGGTGCCTGACCCGATATTACTCAAGCTTCATTAATCTGGAGAGAGATC 2100

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           |||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||
Db    2101 TAGCTTCAGGACTCATTGGCCCTCTTCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 2160

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||||||||| |||||||||||||| |||||||||||||||||||||||||||||||| |
Db    2161 GAAACCAGATGATGTCAGACAAGAGAAATGTCATCCTGTTTTCTGTATTTGATGAGAATC 2220

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           |||||||||||||||||||||||| || ||||| ||||||||| ||| || ||||||||
Db    2221 GAAGCTGGTACCTCACAGAGAATATGCAGCGCTTCCTCCCCAATGCAGATGTAGTGCAGC 2280

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
              | || ||||||||||||  ||| ||||||||||||||||||||||||||||||||||
Db    2281 CCCATGACCCAGAGTTCCAACTCTCTAACATCATGCACAGCATCAATGGCTATGTTTTTG 2340

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           | |  |||||| |||||||||||||||||||||||| ||||||||||||||||  |||
Db    2341 ACAACTTGCAGCTGTCAGTTTGTTTGCATGAGGTGGCGTACTGGTACATTCTAAGTGTTG 2400

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           ||||||| |||||||||| |||||||||||||||||||||||||||||||||||||||||
Db    2401 GAGCACAAACTGACTTCCTGTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2460

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           |||||||||||||||| ||||| ||||||||||||||||||||||||||||||| ||||
Db    2461 TCTATGAAGACACACTTACCCTCTTCCCATTCTCAGGAGAAACTGTCTTCATGTCAATGG 2520

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           |||||||||||||| ||| ||||||||||||||||||||||||||||||||||||||||
Db    2521 AAAACCCAGGTCTGTGGGTTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2580

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           | |||||||||||||||||||||||| ||| ||||| || ||||||||| ||||||| |
Db    2581 CAGCCTTACTGAAGGTTTCTAGTTGTAACAGGAACATTGATGATTATTATGAGGACACAT 2640

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           | ||||||||| || |   | |||| | | |||||||||  ||| |||| ||||||||||
Db    2641 ACGAAGATATTCCAACTCCCCTGCTAAATGAAAACAATGTAATTAAACCTAGAAGCTTCT 2700

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
           |||||||| ||     |   |       |  ||    || | |
Db    2701 CCCAGAATTCAAGGCACCCTAGCACTAAGGAAAAGCAATTGA---------------- 2742

Qy     721 CAGATCAAGAGGAAATTGACTATGATGATACCATATCAGTTGAAATGAAGAAGGAAGATT 780
                                                    |||||||| |||||||
Db    2743 ----------------------------------------AAATGAAGAGAGAAGATT 2760

Qy     781 TTGACATTTATGATGAGGATGAAAATCAGAGCCCCCGCAGCTTTCAAAAGAAAACACGAC 840
           ||||||| || |  ||  |||||||||||| ||| ||||||||||||||||||||||||
Db    2761 TTGACATCTACGGCGACTATGAAAATCAGGGCCTCCGCAGCTTTCAAAAGAAAACACGAC 2820

Qy     841 ACTATTTTATTGCTGCAGTGGAGAGGCTCTGGGATTATGGGATGAGTAGCTCCCCACATG 900
           ||||||| |||||||||||||| | |||||||||||||||||||||||| || || |||
Db    2821 ACTATTTCATTGCTGCAGTGGAGCGTCTCTGGGATTATGGGATGAGTAGATCTCCCCATA 2880

Qy     901 TTCTAAGAAACAGGGCTCAGAGTGGCAGTGTCCCTCAGTTCAAGAAAGTTGTTTTCCAGG 960
           | |||||||||||||||| |||||   |||||  |||||||||| || ||||||||||
Db    2881 TACTAAGAAACAGGGCTCAAAGTGGGGATGTCCAGCAGTTCAAGAAGGTGGTTTTCCAGG 2940

Qy     961 AATTTACTGATGGCTCCTTTACTCAGCCCTTATACCGTGGA 1001
           ||||||||||||| ||||||||||||||||||||||||||||
Db    2941 AATTTACTGATGGATCCTTTACTCAGCCCTTATACCGTGGA 2981

RESULT 8
US-09-957-641-1
; Sequence 1, Application US/09957641
; Publication No. US20020182670A1
; GENERAL INFORMATION:
; APPLICANT: Emory University
; TITLE OF INVENTION: MODIFIED FACTOR VIII
; FILE REFERENCE: 75-00
; CURRENT APPLICATION NUMBER: US/09/957,641
; CURRENT FILING DATE: 2001-09-16
; PRIOR APPLICATION NUMBER: US 60/234047
; PRIOR FILING DATE: 2000-09-19
; PRIOR APPLICATION NUMBER: US 60/236460
; PRIOR FILING DATE: 2000-09-29
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
; LENGTH: 9009
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (208)..(7203)
US-09-957-641-1

Query Match 67.0%; Score 671; DB 9; Length 9009;
Best Local Similarity 94.6%; Pred. No. 6e-188;
Matches 695; Conservative 0; Mismatches 40; Indels 0; Gaps 0;

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Qy   1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1775 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 1834
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Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1835 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 1894

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1895 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 1954

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1955 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 2014

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2015 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 2074

Qy     301 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2075 ATAGTTTGCAGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTG 2134

Qy     361 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2135 GAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGG 2194

Qy     421 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2195 TCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGG 2254

Qy     481 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2255 AAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGA 2314

Qy     541 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2315 CCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACAGTT 2374

Qy     601 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2375 ATGAAGATATTTCAGCATACTTGCTGAGTAAAAACAATGCCATTGAACCAAGAAGCTTCT 2434

Qy     661 CCCAGAATCCACCAGTCTTGAAACGCCATCAACGCGAAATAACTCGTACTACTCTTCAGT 720
           |||||||| ||  |  |   |         |||    || | | |   || ||  |||
Db    2435 CCCAGAATTCAAGACACCCTAGCACTAGGCAAAAGCAATTTAATGCCACCACAATTCCAG 2494

Qy     721 CAGATCAAGAGGAAA 735
            | || |    || |
Db    2495 AAAATGACATAGAGA 2509


RESULT 9
US-09-880-107-2275
; Sequence 2275, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,107
;  CURRENT FILING DATE:  2001-06-14
;  PRIOR APPLICATION NUMBER: US 60/211,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2275
;   LENGTH: 6909
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020142981A1 M16967
US-09-880-107-2275

  Query Match             15.3%;  Score 153.6;  DB 10;  Length 6909;
  Best Local Similarity   54.4%;  Pred. No. 4.6e-35;
  Matches  331;  Conservative    0;  Mismatches  274;  Indels    3;  Gaps    1;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||   ||| | | |||| | ||  |    ||||  |||  |||  | ||   ||||||
Db    1574 AAAATGATGCCCAGTGCTTAACAAGACCATACTACAGTGACGTGGACATCATGAGAGACA 1633

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | || || || || || || | || || |||||  | |    |||  | ||     |||
Db    1634 TCGCCTCTGGGCTAATAGGACTACTTCTAATCTGTAAGAGCAGATCCCTGGACAGGCGAG 1693

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||  ||||      ||||||      |         ||||| |||| ||||||||||||
Db    1694 GAATACAGAGGGCAGCAGACATCGAACAGCAGGCTGTGTTTGCTGTGTTTGATGAGAACA 1753

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||||    || || ||  |    |||      ||||| | ||  ||| | |
Db    1754 AAAGCTGGTACCTTGAGGACAACATCAACAAGTTTTGTGAAAATCCTGATGAGGTGAAAC 1813

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           |||  || ||  ||||  | |  || |||||||||  ||  ||||||||||||||   ||
Db    1814 GTGATGACCCCAAGTTTTATGAATCAAACATCATGAGCACTATCAATGGCTATGTGCCTG 1873

Qy     301 ATAGTTTGC---AGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCA 357
           | ||  |         |   | | || ||  ||||    |   | ||| || |    ||
Db    1874 AGAGCATAACTACTCTTGGATTCTGCTTTGATGACACTGTCCAGTGGCACTTCTGTAGTG 1933

Qy     358 TTGGAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAA 417
           | ||  | |||| |||  |  |  |  ||  |||| ||||  |  | ||||   |   ||
Db    1934 TGGGGACCCAGAATGAAATTTTGACCATCCACTTCACTGGGCACTCATTCATCTATGGAA 1993

Qy     418 TGGTCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGA 477
           |   |||| |||||  | |||||  |||||  |    ||||||  |||     |  | |
Db    1994 AGAGGCATGAGGACACCTTGACCCTCTTCCCCATGCGTGGAGAATCTGTGACGGTCACAA 2053

Qy     478 TGGAAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCA 537
           |||| ||   ||    |||||  |    | |   || ||              | | || |
Db    2054 TGGATAATGTTGGAACTTGGATGTTAACTTCCATGAATTCTAGTCCAAGAAGCAAAAAGC 2113

Qy     538 TGACCGCCTTACTGAAGGTTTCTAGTTGTGACAAGAACACTGGTGATTATTACGAGGACA 597
           |||       | | | || |  ||  |||  |    |   || |||  | |   | || |
Db    2114 TGAGGCTGAAATTCAGGGATGTTAAATGTATCCCAGATGATGATGAAGACTCATATGAGA 2173

Qy     598 GTTATGAA 605
            || ||||
Db    2174 TTTTTGAA 2181


RESULT 10
US-09-917-800A-1539
; Sequence 1539, Application US/09917800A
; Patent No. US20020119462A1
; GENERAL INFORMATION:
;  APPLICANT: Mendrick, Donna
;  APPLICANT:  Porter, Mark
;  APPLICANT:  Johnson, Kory
;  APPLICANT:  Castle, Arthur
;  APPLICANT:  Elashoff, Michael
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Molecular Toxicology Modeling
;  FILE REFERENCE: 44921-5038-US
;  CURRENT APPLICATION NUMBER: US/09/917,800A
;  CURRENT FILING DATE:  2001-07-31
;  PRIOR APPLICATION NUMBER: US 60/222,040
;  PRIOR FILING DATE: 2000-07-31
;  PRIOR APPLICATION NUMBER: US 60/222,880
;  PRIOR FILING DATE: 2000-11-02
;  PRIOR APPLICATION NUMBER: US 60/290,029
;  PRIOR FILING DATE: 2001-05-11
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;  PRIOR APPLICATION NUMBER: US 60/290,645
;  PRIOR FILING DATE: 2001-05-15
;  PRIOR APPLICATION NUMBER: US 60/292,336
;  PRIOR FILING DATE: 2001-05-22
;  PRIOR APPLICATION NUMBER: US 60/295,798
;  PRIOR FILING DATE: 2001-06-06
;  PRIOR APPLICATION NUMBER: US 60/297,457
;  PRIOR FILING DATE: 2001-06-13
;  PRIOR APPLICATION NUMBER: US 60/298,884
;  PRIOR FILING DATE: 2001-06-19
;  PRIOR APPLICATION NUMBER: US 60/303,459
;  PRIOR FILING DATE: 2001-07-09
;  NUMBER OF SEQ ID NOS: 1740
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1539
;   LENGTH: 3700
;   TYPE: DNA
;   ORGANISM: Rattus norvegicus
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020119462A1 NM_012532
US-09-917-800A-1539

  Query Match             12.4%;  Score 124.4;  DB 10;  Length 3700;
  Best Local Similarity   53.7%;  Pred. No. 1.4e-26;
  Matches  304;  Conservative    0;  Mismatches  256;  Indels    6;  Gaps    2;

Qy      15 TGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATCTAGCTTCAGGACTC 74
           ||  ||||| |   ||||     |   ||  ||       | |||| | || ||||||||
Db     532 TGTGTGACCAGGATTTACCACTCTCATGTGGATGCTCCAAAAGATATTGCATCAGGACTC 591

Qy      75 ATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAGGAAACCAGATAATG 134
           || || ||  |  |  ||||  | ||||  ||| |  || |    | |||  ||  ||
Db     592 ATAGGACCGTTGATACTCTGTAAAAAAGGTTCTCTGCATAA---GGAAAAAGAGGAAAAT 648

Qy     135 TCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACCGAAGCTGGTACCTC 194
             ||| |       ||  |  |||| |||||  | |||||  || |  ||||||||||
Db     649 ATTGACCAAGAATTCGTACTGATGTTCTCTGTGGTGGATGAAAATCTCAGCTGGTACCTA 708

Qy     195 ACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGCTTGAGGATCCAGAG 254
           ||| ||  ||  ||  |||   | |  | ||||    |||  |    ||  ||  |||
Db     709 GAAGATAACATCAAAACCTTCTGCTCTGAACCAGAGAAAGTCGATAAAGACAATGAAGAC 768

Qy     255 TTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTGATAGTTTGC---AG 311
           |||||  |    |||||  ||| ||   || ||||| |||   ||||  ||  | |    |
Db     769 TTCCAGGAAAGCAACAGGATGTACTCTATAAATGGATATACATTTGGAAGCCTCCCAGGG 828

Qy     312 TTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCATTGGAGCACAGACT 371
           | ||  | |||    | ||    |   | ||||| || |  | || ||      |   |
Db     829 CTCTCGATGTGTGCAGAAGACAGAGTGAAGTGGTACCTTTTTGGGATGGGGAATGAAGTT 888

Qy     372 GACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAATGGTCTATGAAGAC 431
           ||| | | ||| |   ||||   |||  |  || | |    ||| |      ||   ||
Db     889 GACGTGCATTCAGAGCTCTTTCATGGTCAAGCCCTGACCAGCAAGAACTATCATACTGAT 948

Qy     432 ACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGATGGAAAACCCAGGT 491
           | | ||| ||| |||||   | |   |     ||| |  |||  |  | || || ||
Db     949 ATAATCAACCTGTTCCCTGCCACTCTAATTGATGTTTCTATGGTGGCCCAGAATCCTGGA 1008

Qy     492 CTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCATGACCGCCTTACTG 551
           | ||||| ||  | ||||| |||    ||  || |||    ||  ||   |||||  |
Db    1009 GTCTGGATGCTCAGTTGCCAGAACCTGAACCATCTGAAAGCTGGTTTGCAGGCCTTTTTC 1068

Qy     552 AAGGTTTCTAGTTGTGACAAGAACAC 577
           |||||  |   ||  |||||  | |
Db    1069 CAGGTTCGTGACTGCAACAAGCCCTC 1094


RESULT 11
US-09-970-966-175
; Sequence 175, Application US/09970966
; Patent No. US20020173638A1
; GENERAL INFORMATION:
;  APPLICANT: Stolk, John A.
;  APPLICANT:  Molesh, David Alan
;  APPLICANT:  Fling, Steven P.
;  APPLICANT:  Xu, Jiangchun
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.484C6
;  CURRENT APPLICATION NUMBER: US/09/970,966
;  CURRENT FILING DATE:  2001-10-02
;  NUMBER OF SEQ ID NOS: 215
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 175
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-970-966-175

  Query Match             11.7%;  Score 117.4;  DB 9;  Length 3321;
  Best Local Similarity   52.7%;  Pred. No. 1.6e-24;
  Matches  303;  Conservative    0;  Mismatches  266;  Indels    6;  Gaps    2;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||  ||||      ||  ||||  |   ||||  |     | || ||      | ||||
Db     506 AAGGAGATGGCAATTGTGTGACTAGGATTTACCATTCCCACATTGATGCTCCAAAAGATA 565

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | || |||||||||| || ||| |  | |||||  | ||||| ||| ||||| |   ||
Db .   566 TTGCCTCAGGACTCATCGGACCTTTAATAATCTGTAAAAAAGATTCTCTAGATAA---AG 622

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||  | | |      |||         |||  |  |||||||||  | ||||| ||
Db     623 AAAAAGAAAAACATATTGACCGAGAATTTGTGGTGATGTTTTCTGTGGTGGATGAAAATT 682

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||   ||| || ||  ||  ||    | |  | ||||    ||| |
Db     683 TCAGCTGGTACCTAGAAGACAACATTAAAACCTACTGCTCAGAACCAGAGAAAGTTGACA 742

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||  |   ||| ||||| |     ||||  ||| |     | ||||| ||  |||||
Db     743 AAGACAACGAAGACTTCCAGGAGAGTAACAGAATGTATTCTGTGAATGGATACACTTTTG 802

Qy     301 ATAGTTTGC---AGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCA 357
           |||  | |      | ||  | |||    | ||    | | | |||||| || |  | |
Db     803 GAAGTCTCCCAGGACTCTCCATGTGTGCTGAAGACAGAGTAAAATGGTACCTTTTTGGTA 862

Qy     358 TTGGAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAA 417
           | ||     |   |||  | |   | |  ||||   ||  |  | | |   |||| |
Db     863 TGGGTAATGAAGTTGATGTGCACGCAGCTTTCTTTCACGGGCAAGCACTGACTAACAAGA 922

Qy     418 TGGTCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGA 477
           |  |   ||||||  ||| ||| || ||     |          ||  |  |||  |
Db     923 ACTACCGTATTGACACAATCAACCTCTTTCCTGCTACCCTGTTTGATGCTTATATGGTGG 982

Qy     478 TGGAAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCA 537
           | ||||| ||   |||||| ||  | || || ||   | ||  || |||    ||
Db     983 CCCAGAACCCTGGAGAATGGATGCTCAGCTGTCAGAATCTAAACCATCTGAAAGCCGGTT 1042

Qy     538 TGACCGCCTTACTGAAGGTTTCTAGTTGTGACAAG 572
           ||   |||||  |  ||||        ||| |||||
Db    1043 TGCAAGCCTTTTTCCAGGTCCAGGAGTGTAACAAG 1077


RESULT 12
US-09-825-294-175
; Sequence 175, Application US/09825294
; Patent No. US20020044491A1
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Stolk, John A.
;  APPLICANT:  Algate, Paul A.
;  APPLICANT:  Fling, Steven P.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
;  TITLE OF INVENTION:  THERAPY AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.484C5
```

```
;  CURRENT APPLICATION NUMBER: US/09/825,294
;  CURRENT FILING DATE:  2001-04-03
;  NUMBER OF SEQ ID NOS: 215
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 175
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-825-294-175

  Query Match             11.7%;  Score 117.4;  DB 10;  Length 3321;
  Best Local Similarity   52.7%;  Pred. No. 1.6e-24;
  Matches  303;  Conservative    0;  Mismatches  266;  Indels    6;  Gaps    2;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||  ||||      ||  |||| |   |||| |     | || ||      | ||||
Db     506 AAGGAGATGGCAATTGTGTGACTAGGATTTACCATTCCCACATTGATGCTCCAAAAGATA 565

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | || ||||||||||| || ||| |  | |||||  | ||||| ||| ||||| |   ||
Db     566 TTGCCTCAGGACTCATCGGACCTTTAATAATCTGTAAAAAAGATTCTCTAGATAA---AG 622

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||  | | |      |||         |||  | ||||||||||  | ||||| ||
Db     623 AAAAAGAAAAACATATTGACCGAGAATTTGTGGTGATGTTTTCTGTGGTGGATGAAAATT 682

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||   ||| || ||  ||  ||    | |  | ||||    |||  |
Db     683 TCAGCTGGTACCTAGAAGACAACATTAAAACCTACTGCTCAGAACCAGAGAAAGTTGACA 742

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||  |   ||| ||||| |     ||||  ||| |     | ||||| ||   |||||
Db     743 AAGACAACGAAGACTTCCAGGAGAGTAACAGAATGTATTCTGTGAATGGATACACTTTTG 802

Qy     301 ATAGTTTGC---AGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCA 357
           |||  | |      | ||  | |||   | ||    | | | |||||| || |  | |
Db     803 GAAGTCTCCCAGGACTCTCCATGTGTGCTGAAGACAGAGTAAAATGGTACCTTTTTGGTA 862

Qy     358 TTGGAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAA 417
           | ||     |   |||  | |   | |  |||||    ||  |  |  | |   |||| |
Db     863 TGGGTAATGAAGTTGATGTGCACGCAGCTTTCTTTCACGGGCAAGCACTGACTAACAAGA 922

Qy     418 TGGTCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGA 477
           |  |   |||||| ||| ||| || ||     |          ||  |  |||  |
Db     923 ACTACCGTATTGACACAATCAACCTCTTTCCTGCTACCCTGTTTGATGCTTATATGGTGG 982

Qy     478 TGGAAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCA 537
           | ||||| ||   |||||| ||  | || || ||   | ||  || |||    ||
Db     983 CCCAGAACCCTGGAGAATGGATGCTCAGCTGTCAGAATCTAAACCATCTGAAAGCCGGTT 1042

Qy     538 TGACCGCCTTACTGAAGGTTTCTAGTTGTGACAAG 572
           ||   |||||  |  ||||       ||| |||||
Db    1043 TGCAAGCCTTTTTCCAGGTCCAGGAGTGTAACAAG 1077


RESULT 13
US-09-880-107-2253
; Sequence 2253, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,107
;  CURRENT FILING DATE:  2001-06-14
;  PRIOR APPLICATION NUMBER: US 60/211,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2253
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020142981A1 M13699
US-09-880-107-2253

  Query Match             11.7%;  Score 117.4;  DB 10;  Length 3321;
  Best Local Similarity   52.7%;  Pred. No. 1.6e-24;
  Matches  303;  Conservative    0;  Mismatches  266;  Indels    6;  Gaps    2;

Qy       1 AATCAGATCCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATC 60
           ||  ||||      ||  |||| |   |||| |     | || ||      | ||||
Db     506 AAGGAGATGGCAATTGTGTGACTAGGATTTACCATTCCCACATTGATGCTCCAAAAGATA 565

Qy      61 TAGCTTCAGGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAG 120
           | || ||||||||||| || ||| |  | |||||  | ||||| ||| ||||| |   ||
Db     566 TTGCCTCAGGACTCATCGGACCTTTAATAATCTGTAAAAAAGATTCTCTAGATAA---AG 622

Qy     121 GAAACCAGATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACC 180
           |||  | | |      |||         |||  | ||||||||||  | ||||| ||
Db     623 AAAAAGAAAAACATATTGACCGAGAATTTGTGGTGATGTTTTCTGTGGTGGATGAAAATT 682

Qy     181 GAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGC 240
           ||||||||||   ||| || ||  ||  ||    | |  | ||||    |||  |
Db     683 TCAGCTGGTACCTAGAAGACAACATTAAAACCTACTGCTCAGAACCAGAGAAAGTTGACA 742

Qy     241 TTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTG 300
           ||  |   ||| ||||| |     ||||  ||| |     | ||||| ||   |||||
Db     743 AAGACAACGAAGACTTCCAGGAGAGTAACAGAATGTATTCTGTGAATGGATACACTTTTG 802

Qy     301 ATAGTTTGC---AGTTGTCAGTTTGTTTGCATGAGGTGGCATACTGGTACATTCTAAGCA 357
           |||  | |      | ||  | |||   | ||    | | | |||||| || |  | |
Db     803 GAAGTCTCCCAGGACTCTCCATGTGTGCTGAAGACAGAGTAAAATGGTACCTTTTTGGTA 862

Qy     358 TTGGAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGGATATACCTTCAAACACAAAA 417
           | ||     |   |||  | |   | |  |||||    ||  |  |  | |   |||| |
Db     863 TGGGTAATGAAGTTGATGTGCACGCAGCTTTCTTTCACGGGCAAGCACTGACTAACAAGA 922

Qy     418 TGGTCTATGAAGACACACTCACCCTATTCCCATTCTCAGGAGAAACTGTCTTCATGTCGA 477
           |  |   |||||| ||| ||| || ||     |          ||  |  |||  |
Db     923 ACTACCGTATTGACACAATCAACCTCTTTCCTGCTACCCTGTTTGATGCTTATATGGTGG 982

Qy     478 TGGAAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTCAGACTTTCGGAACAGAGGCA 537
           | ||||| ||   |||||| ||  | || || ||   | ||  || |||    ||
Db     983 CCCAGAACCCTGGAGAATGGATGCTCAGCTGTCAGAATCTAAACCATCTGAAAGCCGGTT 1042

Qy     538 TGACCGCCTTACTGAAGGTTTCTAGTTGTGACAAG 572
           ||   |||||  |  ||||       ||| |||||
Db    1043 TGCAAGCCTTTTTCCAGGTCCAGGAGTGTAACAAG 1077


RESULT 14
US-09-864-864-114
; Sequence 114, Application US/09864864
; Patent No. US20020102679A1
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Mitcham, Jennifer L.
;  APPLICANT:  Harlocker, Susan L.
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Secrist, Heather
;  APPLICANT:  Lodes, Michael J.
;  APPLICANT:  Algate, Paul A.
;  APPLICANT:  Fling, Steve P.
;  APPLICANT:  Mannion, Jane
;  APPLICANT:  Benson, Darin R.
;  APPLICANT:  Carter, Darrick
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.523
```

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;  CURRENT APPLICATION NUMBER: US/09/864,864
;  CURRENT FILING DATE:  2001-05-23
;  NUMBER OF SEQ ID NOS: 341
;  SOFTWARE: Corixa Invention Disclosure Database
; SEQ ID NO 114
;   LENGTH: 596
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(596)
;   OTHER INFORMATION: n = A,T,C or G
US-09-864-864-114

  Query Match             6.3%;  Score 63;  DB 10;  Length 596;
  Best Local Similarity   49.6%;  Pred. No. 7.8e-09;
  Matches  190;  Conservative    0;  Mismatches  190;  Indels    3;  Gaps    1;

Qy    161 TTCTGTATTTGATGAGAACCGAAGCTGGTACCTCACAGAGAATATACAACGCTTTCTCCC 220
          | | ||||||||||||||    || |   |||   ||| |||||   |   |||    |
Db     12 TACAGTATTTGATGAGAATGAGAGTTTACTCCTGGAAGATAATATTAGAATGTTTACAAC 71

Qy    221 CAATCCAGCTGGAGTGCAGCTTGAGGATCCAGAGTTCCAAGCCTCCAACATCATGCACAG 280
            || | |   ||| |    || |||  ||| || || |  || || |  ||||||
Db     72 TGCACCTGATCAGGTGGATAAGGAAGATGAAGACTTTCAGGAATCTAATAAAATGCACTC 131

Qy    281 CATCAATGGCTATGTTTTTGATAGTTTGCAGT---TGTCAGTTTGTTTGCATGAGGTGGC 337
          ||| |||||  |   | | ||  | |  || |    |  |  | ||       ||   ||
Db    132 CATGAATGGATTCATGTATGGGAATCAGCCGGGTCTCACTATGTGCAAAGGAGATTCGGT 191

Qy    338 ATACTGGTACATTCTAAGCATTGGAGCACAGACTGACTTCCTTTCTGTCTTCTTCTCTGG 397
            |||||| |  | |||   |||    || | ||  | | |    | | ||| || ||
Db    192 CGTGTGGTACTTATTCAGCGCCGGAAATGAGGCCGATGTACATGGAATATACTTTTCAGG 251

Qy    398 ATATACCTTCAAACACAAAATGGTCTATGAAGACACACTCACCCTATTCCCATTCTCAGG 457
          | | || |        | |   |       |||||||   | ||| |||||     || |
Db    252 AAACACATATCTGTGGAGAGGAGAACGGAGAGACACAGCAAACCTCTTCCCTCAAACAAG 311

Qy    458 AGAAACTGTCTTCATGTCGATGGAAAACCCAGGTCTATGGATTCTGGGGTGCCACAACTC 517
            ||  ||  ||||| |   || |     ||    |  | | | |  ||||  |   |
Db    312 TCTTACGCTCCACATGTGGCCTGACACAGAGGGGACTTTTAATGTTGAATGCCTTACAAC 371

Qy    518 AGACTTTCGGAACAGAGGCATGA 540
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Db    372 TGATCATTACACAGGCGGCATGA 394


RESULT 15
US-09-778-320-132/c
; Sequence 132, Application US/09778320
; Patent No. US20010034052A1
; GENERAL INFORMATION:
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Day, Craig H.
;  APPLICANT:  Jiang, Yuqiu
;  APPLICANT:  Houghton, Raymond L.
;  APPLICANT:  Mitcham, Jennifer
;  APPLICANT:  Wang, TongTong
;  APPLICANT:  McNeill, Patricia D.
;  TITLE OF INVENTION:  COMPOSITIONS AND METHODS FOR THE THERAPY AND
;  TITLE OF INVENTION:  DIAGNOSIS OF BREAST CANCER
;  FILE REFERENCE:  210121.491C5
;  CURRENT APPLICATION NUMBER: US/09/778,320
;  CURRENT FILING DATE:  2001-02-06
;  NUMBER OF SEQ ID NOS:  301
;  SOFTWARE:  FastSEQ for Windows Version 3.0
; SEQ ID NO 132
;   LENGTH: 404
;   TYPE: DNA
;   ORGANISM: Homo sapien
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(404)
;   OTHER INFORMATION: n = A,T,C or G
US-09-778-320-132

  Query Match             6.2%;  Score 61.8;  DB 10;  Length 404;
  Best Local Similarity   52.0%;  Pred. No. 1.4e-08;
  Matches  158;  Conservative    0;  Mismatches  145;  Indels    1;  Gaps    1;

Qy      9 CCTCGGTGCCTGACCCGCTATTACTCTAGTTTCGTTAATATGGAGAGAGATCTAGCTTCA 68
          |||  ||| ||  |        || | |  |   ||  ||      | |||| ||    |
Db    341 CCTGTGTGTCTAGCTAAGATGTATTATTCTGCTGTGGATCCCACTAAAGATATATTCACT 282

Qy     69 GGACTCATTGGCCCTCTCCTCATCTGCTACAAAGAATCTGTAGATCAAAGAGGAAACCAG 128
          || || ||||| ||  |    || ||| | |||| |  | || ||  ||  || | |||
Db    281 GGGCTTATTGGGCCAATGAAAATATGCAAGAAAGGAAGTTTATATGCAAATGGGAGACAG 222

Qy    129 ATAATGTCAGACAAGAGGAATGTCATCCTGTTTTCTGTATTTGATGAGAACCGAAGCTGG 188
          | |     |||||||          |  |   | | |||||||||||||||    || |
Db    221 AAAGATGTAGACAAGGAATTCTATTTGTTTCCTACAGTATTTGATGAGAATGAGAGTTTA 162

Qy    189 TACCTCACAGAGAATA-TACAACGCTTTCTCCCCAATCCAGCTGGAGTGCAGCTTGAGGA 247
            |||   ||| |||| ||       |||    |    || | |   ||| |    || ||
Db    161 CTCCTGGAAGATAATATTAGGNATGTTTACAACTGCNCCTGATCAGGTGGATAAGGAAGA 102

Qy    248 TCCAGAGTTCCAAGCCTCCAACATCATGCACAGCATCAATGGCTATGTTTTTGATAGTTT 307
          |  ||| || || |  || || |  ||||||  ||| ||||| |   | | ||  | |
Db    101 TGAAGACTTTCAGGNATCTAATAAAATGCACTCCATGAATGGATTCATGTATGGGAATCA 42

Qy    308 GCAG 311
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Db     41 GCCG 38


Search completed: January  5, 2003, 03:12:15
Job time : 118 secs
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GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: January 4, 2003, 23:31:50 ; Search time 47 Seconds
(without alignments)
6531.563 Million cell updates/sec

Title: US-09-740-211-14_COPY_3600_4600
Perfect score: 1001
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1001	100.0	4629	2	US-08-484-891-7	Sequence 7, Appli
2	1001	100.0	4670	3	US-08-717-294-41	Sequence 41, Appl
3	1001	100.0	4999	4	US-09-470-618-14	Sequence 14, Appl
4	1001	100.0	4999	4	US-09-364-862-14	Sequence 14, Appl
5	1001	100.0	5035	2	US-08-882-083-1	Sequence 1, Appli
6	1001	100.0	5035	2	US-08-558-107-1	Sequence 1, Appli
7	1001	100.0	5035	3	US-09-243-539-1	Sequence 1, Appli
8	1001	100.0	7056	1	US-08-121-202-1	Sequence 1, Appli
9	1001	100.0	8967	1	US-08-366-851A-1	Sequence 1, Appli
10	1001	100.0	9009	1	US-07-864-004B-3	Sequence 3, Appli
11	1001	100.0	9009	1	US-08-251-937A-3	Sequence 3, Appli
12	1001	100.0	9009	1	US-08-212-133A-1	Sequence 1, Appli
13	1001	100.0	9009	1	US-08-474-503-1	Sequence 1, Appli
14	1001	100.0	9009	2	US-08-670-707A-1	Sequence 1, Appli
15	1001	100.0	9009	4	US-09-037-601-1	Sequence 1, Appli
16	1001	100.0	9009	4	US-09-315-179-1	Sequence 1, Appli
17	1001	100.0	9009	4	US-09-523-656-1	Sequence 1, Appli
18	1001	100.0	9009	5	PCT-US93-03275-3	Sequence 3, Appli
19	1001	100.0	9009	5	PCT-US94-13200-1	Sequence 1, Appli
20	1001	100.0	9354	1	US-08-683-839B-2	Sequence 2, Appli
21	1001	100.0	11933	4	US-09-470-618-13	Sequence 13, Appl
22	1001	100.0	11933	4	US-09-364-862-13	Sequence 13, Appl
23	999.4	99.8	8241	6	5171844-1	Patent No. 5171844
24	997.8	99.7	6999	1	US-08-276-594A-1	Sequence 1, Appli
25	843.2	84.2	7493	1	US-08-212-133A-7	Sequence 7, Appli
26	843.2	84.2	7493	1	US-08-474-503-5	Sequence 5, Appli
27	843.2	84.2	7493	2	US-08-670-707A-5	Sequence 5, Appli
28	843.2	84.2	7493	4	US-09-037-601-5	Sequence 5, Appli
29	843.2	84.2	7493	4	US-09-315-179-5	Sequence 5, Appli
30	843.2	84.2	7493	5	PCT-US94-13200-5	Sequence 5, Appli
31	841.6	84.1	7032	4	US-09-324-867-1	Sequence 1, Appli
32	808.8	80.8	4334	2	US-08-670-707A-38	Sequence 38, Appl
33	808.8	80.8	4334	4	US-09-037-601-38	Sequence 38, Appl
34	808.8	80.8	4334	4	US-09-315-179-38	Sequence 38, Appl
35	808.8	80.8	4404	4	US-09-523-656-37	Sequence 37, Appl
36	808.8	80.8	6402	2	US-08-670-707A-36	Sequence 36, Appl
37	808.8	80.8	6402	4	US-09-037-601-36	Sequence 36, Appl
38	808.8	80.8	6402	4	US-09-315-179-36	Sequence 36, Appl
39	808.8	80.8	6402	4	US-09-523-656-29	Sequence 29, Appl
40	609	60.8	4451	3	US-08-717-294-42	Sequence 42, Appl
41	202.2	20.2	6585	3	US-08-746-111-4	Sequence 4, Appli
42	198.2	19.8	6909	2	US-08-804-196-1	Sequence 1, Appli
43	198.2	19.8	6909	2	US-08-658-340-1	Sequence 1, Appli
44	198.2	19.8	6909	3	US-08-746-111-26	Sequence 26, Appl
45	93.4	9.3	1780	2	US-08-480-229C-11	Sequence 11, Appl

ALIGNMENTS

RESULT 1
US-08-484-891-7
; Sequence 7, Application US/08484891
; Patent No. 5935935
; GENERAL INFORMATION:
; APPLICANT: Connelly, Sheila
; APPLICANT: Kaleko, Michael
; APPLICANT: Smith, Theodore
; TITLE OF INVENTION: Adenoviral Vectors for
; TITLE OF INVENTION: Treatment of Hemophilia
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carella, Byrne, Bain, Gilfillan,
; ADDRESSEE: Cecchi, Stewart & Olstein
; STREET: 6 Becker Farm Road
; CITY: Roseland
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 inch diskette
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WordPerfect 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/484,891
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/218,335
; FILING DATE: 25-MAR-1994
; APPLICATION NUMBER: 08/074,920
; FILING DATE: 10-JUN-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Olstein, Elliot M.
; REGISTRATION NUMBER: 24,025
; REFERENCE/DOCKET NUMBER: 271010-273
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 201-994-1700
; TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4629 bases
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA primer
; FEATURE:
; NAME/KEY: Factor VIII cDNA with
; NAME/KEY: B domain deleted

US-08-484-891-7

Query Match 100.0%; Score 1001; DB 2; Length 4629;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3227 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3286

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3287 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3346

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3347 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3406

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3407 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3466

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3467 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3526

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3527 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3586

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3587 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 3646

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3647 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 3706

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3707 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 3766

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3767 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 3826

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3827 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 3886

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3887 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 3946

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3947 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4006

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4007 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4066

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4067 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4126

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4127 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4186

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4187 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4227
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RESULT 2
US-08-717-294-41
; Sequence 41, Application US/08717294
; Patent No. 6114148
; GENERAL INFORMATION:
; APPLICANT: SEED, BRIAN
; APPLICANT: HAAS, JURGEN
; TITLE OF INVENTION: HIGH LEVEL EXPRESSION OF
; TITLE OF INVENTION: PROTEINS
; NUMBER OF SEQUENCES: 110
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Clark & Elbing LLP
; STREET: 176 Federal Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/717,294
; FILING DATE: 20-SEP-1996
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Elbing, Karen L
; REGISTRATION NUMBER: 35,238
; REFERENCE/DOCKET NUMBER: 00786/345001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-428-0200
; TELEFAX: 617-428-7045
; TELEX:
; INFORMATION FOR SEQ ID NO: 41:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4670 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-717-294-41

Query Match 100.0%; Score 1001; DB 3; Length 4670;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3253 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3312

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3313 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3372

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3373 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3432

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3433 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3492

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3493 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3552

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3553 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3612

Qy    361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3613 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 3672

Qy    421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3673 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 3732

Qy    481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3733 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 3792

Qy    541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3793 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 3852

Qy    601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3853 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 3912

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3913 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 3972

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3973 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4032

Qy    781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4033 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4092

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4093 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4152

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4153 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4212

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   4213 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4253
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RESULT 3
US-09-470-618-14
; Sequence 14, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/09/470,618
; CURRENT FILING DATE: 1999-12-22
; EARLIER APPLICATION NUMBER: 09/364,862
; EARLIER FILING DATE: 1999-07-30
; EARLIER APPLICATION NUMBER: 60/125,974
; EARLIER FILING DATE: 1999-03-24
; EARLIER APPLICATION NUMBER: 60/104,994
; EARLIER FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 4999
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-470-618-14

Query Match 100.0%; Score 1001; DB 4; Length 4999;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3600 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3659

Qy     61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3660 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3719

Qy    121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3720 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3779

Qy    181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3780 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3839

Qy    241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3840 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3899

Qy    301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3900 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3959

Qy    361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3960 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4019

Qy    421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4020 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4079

Qy    481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4080 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4139

Qy    541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4140 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4199

Qy    601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4200 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4259

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4260 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4319

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4320 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4379

Qy    781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4380 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4439

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4440 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4499

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4500 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4559

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
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Db   4560 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4600


RESULT 4
US-09-364-862-14
; Sequence 14, Application US/09364862
; Patent No. 6221349
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: ADENO-ASSOCIATED VECTORS FOR EXPRESSION OF FACTOR VIII
;  TITLE OF INVENTION: BY TARGET
;  TITLE OF INVENTION: CELLS
;  FILE REFERENCE: AVIGEN-03743
;  CURRENT APPLICATION NUMBER: US/09/364,862
;  CURRENT FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 14
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-364-862-14

  Query Match             100.0%;  Score 1001;  DB 4;  Length 4999;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3600 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3659

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3660 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3719

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3720 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3779

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3780 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3839

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3840 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3899

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3900 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3959

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3960 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4019

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4020 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4079

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4080 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4139

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4140 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4199

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4200 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4259

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4260 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4319

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4320 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4379

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4380 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4439

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4440 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4499

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4500 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4559

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4560 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4600


RESULT 5
US-08-882-083-1
; Sequence 1, Application US/08882083
; Patent No. 5869292
; GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/882,083
;      FILING DATE:
;      CLASSIFICATION:  514
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
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;      LOCATION:  35..5017
US-08-882-083-1

  Query Match             100.0%;  Score 1001;  DB 2;  Length 5035;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3831 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3890

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3891 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3950

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3951 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 4010

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4011 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 4070

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4071 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 4130

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4131 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 4190

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4191 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4250

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4251 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4310

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4311 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4370

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4371 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4430

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4431 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4490

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4491 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4550

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4551 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4610

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4611 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4670

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4671 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4730

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4731 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4790

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4791 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4831


RESULT 6
US-08-558-107-1
; Sequence 1, Application US/08558107
; Patent No. 5910481
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/558,107
;      FILING DATE:  13-NOV-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  35..5017
US-08-558-107-1

  Query Match             100.0%;  Score 1001;  DB 2;  Length 5035;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3831 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3890

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3891 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3950

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3951 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 4010

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4011 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 4070

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4071 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 4130

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4131 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 4190
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Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4191 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4250

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4251 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4310

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4311 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4370

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4371 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4430

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4431 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4490

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4491 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4550

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4551 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4610

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4611 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4670

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4671 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4730

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4731 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4790

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
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Db    4791 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4831
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RESULT 7
US-09-243-539-1
; Sequence 1, Application US/09243539
; Patent No. 6130203
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/243,539
;      FILING DATE:
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  35..5017
US-09-243-539-1

  Query Match             100.0%;  Score 1001;  DB 3;  Length 5035;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3831 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3890

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3891 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3950

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3951 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 4010

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4011 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 4070

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4071 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 4130

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4131 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 4190

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4191 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4250

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4251 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4310

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4311 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4370

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4371 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4430

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4431 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4490

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db    4491 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4550

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
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Db   4551 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4610

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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4611 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4670

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4671 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4730

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4731 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4790

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   4791 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4831


RESULT 8
US-08-121-202-1
; Sequence 1, Application US/08121202
; Patent No. 5563045
;  GENERAL INFORMATION:
;    APPLICANT:  Pittman, Debra
;    APPLICANT:  Rehemtulla, Alnawaz
;    APPLICANT:  Wozney, John M.
;    APPLICANT:  Kaufman, Randal J.
;    TITLE OF INVENTION:  CHIMERIC PROCOAGULANT PROTEINS
;    NUMBER OF SEQUENCES:  27
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Legal Affairs, Genetics Institute, Inc.
;      STREET:  87 CambridgePark Drive
;      CITY:  Cambridge
;      STATE:  MA
;      COUNTRY:  USA
;      ZIP:  02140
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/121,202
;      FILING DATE:  14-SEP-1993
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Meinert, M. C.
;      REGISTRATION NUMBER:  31,544
;      REFERENCE/DOCKET NUMBER:  GI 5195A
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)876-1210 X8574
;      TELEFAX:  (617) 876-5851
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  7056 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..7053
US-08-121-202-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 7056;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db   5867 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 5926

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5927 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 5986

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5987 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6046

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6047 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6106

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6107 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6166

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6167 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6226

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6227 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6286

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6287 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6346

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6347 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6406

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6407 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6466

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6467 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6526

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6527 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6586

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6587 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6646

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6647 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6706

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6707 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 6766

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6767 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6826

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   6827 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 6867


RESULT 9
US-08-366-851A-1
; Sequence 1, Application US/08366851A
; Patent No. 5681746
;  GENERAL INFORMATION:
;    APPLICANT:  Bodner, Mordechai
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;    APPLICANT:  De Polo, Nicolas J.
;    APPLICANT:  Hsu, David Chi-Tang
;    APPLICANT:  Chang, Steven
;    TITLE OF INVENTION:  Retroviral Delivery of Full Length Factor VIII
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Viagene, Inc.
;      STREET:  11055 Roselle Street
;      CITY:  San Diego
;      STATE:  California
;      COUNTRY:  U.S.A.
;      ZIP:  92121
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/366,851A
;      FILING DATE:
;      CLASSIFICATION:  514
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Chambers, Daniel M.
;      REGISTRATION NUMBER:  34,561
;      REFERENCE/DOCKET NUMBER:  930049.438
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (619) 452-1288
;      TELEFAX:  (619) 452-2616
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  8967 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  110..7165
US-08-366-851A-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 8967;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5976 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6035

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6036 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 6095

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6096 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6155

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6156 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6215

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6216 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6275

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6276 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6335

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6336 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6395

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6396 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6455

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6456 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6515

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6516 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6575

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6576 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6635

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db    6636 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6695

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6696 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6755

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6756 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6815

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6816 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 6875

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6876 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6935

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
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Db    6936 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 6976


RESULT 10
US-07-864-004B-3
; Sequence 3, Application US/07864004B
; Patent No. 5364771
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  6
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/864,004B
;      FILING DATE:  07 APRIL 1992
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU106
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6508
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  3:
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;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5001 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2
;      OTHER INFORMATION:  domain"
US-07-864-004B-3

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6077 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 6136

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6197 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6256

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6437 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6496

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6797 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6856

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6857 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 6916

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
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Db    6917 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6976

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
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Db    6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017


RESULT 11
US-08-251-937A-3
; Sequence 3, Application US/08251937A
; Patent No. 5583209
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  10
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/251,937A
;      FILING DATE:  31-MAY-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106DIV
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6367
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
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;       NAME/KEY:  misc_feature (Domain Structure)
;       LOCATION:  5001 . . . 7053
;       OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;       OTHER INFORMATION:  domain"
;     FEATURE:
;       NAME/KEY:  misc_feature (Domain Structure)
;       LOCATION:  1 . . . 2277
;       OTHER INFORMATION:  /note= "Equivalent to the A1-A2
;       OTHER INFORMATION:  domain"
US-08-251-937A-3

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
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Db    6077 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 6136

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
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Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
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Db    6197 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6256

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
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Db    6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6437 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6496

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
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Db    6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db    6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
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Db    6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
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Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
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Db    6857 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 6916

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
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Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
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Db    6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017


RESULT 12
US-08-212-133A-1
; Sequence 1, Application US/08212133A
; Patent No. 5663060
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/212,133A
;      FILING DATE:  March 11, 1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU/76677
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-572-6508
;      TELEFAX:  404-572-6555
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5125 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2 domain."
;    FEATURE:
;      NAME/KEY:  Domain
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /note= "cDNA encoding human factor
;      OTHER INFORMATION:  VIII."
US-08-212-133A-1
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  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
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Db    6077 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 6136

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6197 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6256

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
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Db    6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
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Db    6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
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Db    6437 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6496

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
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Db    6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
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Db    6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db    6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6797 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6856

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Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
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Db    6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017
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RESULT 13
US-08-474-503-1
; Sequence 1, Application US/08474503
; Patent No. 5744446
;  GENERAL INFORMATION:
;    APPLICANT:  Emory University
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street, Suite 2800
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/474,503
;      FILING DATE:  07-JUN-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106CIP(3)
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6500
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5125 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2 domain."
;    FEATURE:
;      NAME/KEY:  Domain
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /note= "cDNA encoding human factor
;      OTHER INFORMATION:  VIII."
US-08-474-503-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
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Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
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Db    6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
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Db    6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
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Db    6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
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Db    6437 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6496

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
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Db    6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
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Db    6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db    6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
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Db    6797 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6856

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
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Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
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Db    6917 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6976

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
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Db    6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017


RESULT 14
US-08-670-707A-1
; Sequence 1, Application US/08670707A
; Patent No. 5859204
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/670,707A
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Greenlee, Lorance L.
;      REGISTRATION NUMBER:  27,894
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  cDNA to mRNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapiens
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature
;      LOCATION:  5125..7053
;      OTHER INFORMATION:  /product= "Domain Structure"
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2 domain"
;    FEATURE:
;      NAME/KEY:  misc_feature
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /product= "Domain Structure"
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2 domain"
;    FEATURE:
;      NAME/KEY:  misc_feature
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /product= "Domain"
;      OTHER INFORMATION:  /note= "cDNA encoding human factorVIII"
US-08-670-707A-1

  Query Match             100.0%;  Score 1001;  DB 2;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
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Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
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Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
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Qy    241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
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Db   6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy    301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
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Db   6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy    361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
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Db   6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy    421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
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Db   6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy    541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
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Db   6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy    601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
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Db   6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
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Db   6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
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RESULT 15
US-09-037-601-1
; Sequence 1, Application US/09037601
; Patent No. 6180371
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/037,601
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Ferber, Donna M.
;      REGISTRATION NUMBER:  33,878
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  cDNA to mRNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapiens
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature
;      LOCATION:  5125..7053
;      OTHER INFORMATION:  /product= "Domain Structure"
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2 domain"
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;      OTHER INFORMATION:  /note= "cDNA encoding human factorVIII"
US-09-037-601-1

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  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

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Db   6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

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Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
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Db   6917 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6976

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Db   6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017
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Search completed: January 5, 2003, 02:15:46
Job time : 229 secs

GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: January 4, 2003, 23:31:50 ; Search time 54 Seconds
(without alignments)
8017.350 Million cell updates/sec

Title: US-09-740-211-14_COPY_3600_4600
Perfect score: 1001
Sequence: 1 atcaaaggattcgatggtat.....gcagtcaagatggccatcag 1001

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 381593 seqs, 216252194 residues

Total number of hits satisfying chosen parameters: 763186

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*
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2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
	1	1001	100.0	4629	10	US-09-150-811-7	GENERAL INFORMA
	2	1001	100.0	4999	9	US-10-007-968-14	Sequence 14, Appl
	3	1001	100.0	4999	10	US-09-740-211-14	Sequence 14, Appl
	4	1001	100.0	7944	12	US-10-095-718-1	Sequence 1, Appli
	5	1001	100.0	9009	9	US-09-957-641-1	Sequence 1, Appli
	6	1001	100.0	11933	9	US-10-007-968-13	Sequence 13, Appl
	7	1001	100.0	11933	10	US-09-740-211-13	Sequence 13, Appl
	8	841.6	84.1	7914	12	US-10-095-718-3	Sequence 3, Appli
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	10	92.4	9.2	4599	9	US-09-974-298-167	Sequence 167, App
	11	76	7.6	2145	9	US-10-003-132-12	Sequence 12, Appl
	12	72.4	7.2	1270	10	US-09-880-107-3020	Sequence 3020, Ap
	13	69.2	6.9	357	10	US-09-960-352-13850	Sequence 13850, A
	14	61.2	6.1	389	10	US-09-960-352-8113	Sequence 8113, Ap
	15	61	6.1	367	10	US-09-960-352-3109	Sequence 3109, Ap
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	23	57	5.7	3321	9	US-09-970-966-175	Sequence 175, App
	24	57	5.7	3321	10	US-09-825-294-175	Sequence 175, App
	25	57	5.7	3321	10	US-09-880-107-2253	Sequence 2253, Ap
	26	55.6	5.6	5657	9	US-09-974-298-96	Sequence 96, Appl
c	27	52	5.2	379	10	US-09-960-352-12844	Sequence 12844, A
	28	48.2	4.8	3151	9	US-10-003-132-1	Sequence 1, Appli
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	30	42.4	4.2	3700	10	US-09-917-800A-1539	Sequence 1539, Ap
	31	39.4	3.9	476	10	US-09-833-381-1488	Sequence 1488, Ap
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	34	38.6	3.9	45	9	US-09-957-641-18	Sequence 18, Appl
	35	37.6	3.8	519	9	US-09-736-457-1255	Sequence 1255, Ap
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c	42	37.4	3.7	39	9	US-09-957-641-5	Sequence 5, Appli
	43	37.4	3.7	39	9	US-09-957-641-12	Sequence 12, Appl
	44	37.4	3.7	39	9	US-09-957-641-13	Sequence 13, Appl
	45	36.6	3.7	8907	9	US-09-738-626-934	Sequence 934, App

ALIGNMENTS

RESULT 1
US-09-150-811-7
; GENERAL INFORMATION:
; APPLICANT: Connelly, Sheila
; Kaleko, Michael
; Smith, Theodore
; TITLE OF INVENTION: Adenoviral Vectors for Treatment of Hemophilia
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carella, Byrne, Bain, Gilfillan, Cecchi, Stewart & Olste
; STREET: 6 Becker Farm Road
; CITY: Roseland
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 inch diskette
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WordPerfect 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/150,811
; FILING DATE: 10-Sep-1998
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/484,891
; FILING DATE: 07-JUN-1995
; APPLICATION NUMBER: 08/218,335
; FILING DATE: 25-MAR-1994
; APPLICATION NUMBER: 08/074,920
; FILING DATE: 10-JUN-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Olstein, Elliot M.
; REGISTRATION NUMBER: 24,025
; REFERENCE/DOCKET NUMBER: 271010-440
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 973-994-1700
; TELEFAX: 973-994-1744
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-150-811-7

Query Match 100.0%; Score 1001; DB 10; Length 4629;

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Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
           |||||||||||||||||||||||||||||||||||||||||
Db    4187 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4227


RESULT 2
```

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US-10-007-968-14
; Sequence 14, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-14

  Query Match             100.0%;  Score 1001;  DB 9;  Length 4999;
  Best Local Similarity   100.0%;  Pred. No. 4.9e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3600 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3659

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3660 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3719

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3720 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3779

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3780 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3839

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3840 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3899

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3900 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3959

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3960 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4019

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4020 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4079

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4080 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4139

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4140 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4199

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db   4200 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4259

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4260 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4319

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4320 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4379

Qy    781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4380 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4439

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4440 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4499

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4500 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4559

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   4560 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4600


RESULT 3
US-09-740-211-14
; Sequence 14, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-14

  Query Match             100.0%;  Score 1001;  DB 10;  Length 4999;
  Best Local Similarity   100.0%;  Pred. No. 4.9e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3600 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3659

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3660 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3719

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3720 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3779

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3780 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3839

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3840 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3899

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3900 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3959

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3960 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4019

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4020 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4079

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4080 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4139

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4140 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4199

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4200 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4259

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4260 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4319

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4320 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4379

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4380 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4439

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4440 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4499

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4500 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4559

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   4560 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4600


RESULT 4
US-10-095-718-1
; Sequence 1, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
;  PRIOR FILING DATE: 2001-08-22
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;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
;   LENGTH: 7944
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Plasmid pDLZ6 encoding Homo sapiens BDD FVIII
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (420)...(4835)
US-10-095-718-1

  Query Match            100.0%;  Score 1001;  DB 12;  Length 7944;
  Best Local Similarity   100.0%;  Pred. No. 6.3e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3646 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3705

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3706 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3765

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3766 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3825

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3826 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3885

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3886 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3945

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3946 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 4005

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4006 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 4065

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4066 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4125

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4126 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4185

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4186 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4245

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4246 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4305

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4306 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4365

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4366 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4425

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4426 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4485

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4486 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4545

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4546 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4605

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4606 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4646


RESULT 5
US-09-957-641-1
; Sequence 1, Application US/09957641
; Publication No. US20020182670A1
; GENERAL INFORMATION:
;  APPLICANT: Emory University
;  TITLE OF INVENTION: MODIFIED FACTOR VIII
;  FILE REFERENCE: 75-00
;  CURRENT APPLICATION NUMBER: US/09/957,641
;  CURRENT FILING DATE:  2001-09-16
;  PRIOR APPLICATION NUMBER: US 60/234047
;  PRIOR FILING DATE: 2000-09-19
;  PRIOR APPLICATION NUMBER: US 60/236460
;  PRIOR FILING DATE: 2000-09-29
;  NUMBER OF SEQ ID NOS: 18
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 9009
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (208)..(7203)
US-09-957-641-1

  Query Match            100.0%;  Score 1001;  DB 9;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 6.7e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6017 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 6076

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6077 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 6136

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6137 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 6196

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6197 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 6256

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6257 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 6316

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6317 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 6376

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6377 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 6436

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
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Db    6437 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 6496

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6497 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 6556

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6557 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 6616

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6617 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 6676

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6677 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 6736

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6737 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 6796

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6797 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 6856

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6857 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 6916

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6917 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 6976

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    6977 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 7017


RESULT 6
US-10-007-968-13
; Sequence 13, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-13

  Query Match             100.0%;  Score 1001;  DB 9;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 7.8e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3575 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3634

Qy     61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3635 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3694

Qy    121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3695 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3754

Qy    181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3755 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3814

Qy    241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3815 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3874

Qy    301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3875 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3934

Qy    361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3935 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 3994

Qy    421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3995 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4054

Qy    481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4055 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4114

Qy    541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4115 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4174

Qy    601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4175 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4234

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4235 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4294

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4295 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4354

Qy    781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4355 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4414

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4415 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4474

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4475 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4534

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4535 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4575


RESULT 7
US-09-740-211-13
; Sequence 13, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
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;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-13

  Query Match             100.0%;  Score 1001;  DB 10;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 7.8e-295;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3575 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 3634

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3635 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 3694

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3695 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 3754

Qy     181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3755 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 3814

Qy     241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3815 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 3874

Qy     301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3875 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3934

Qy     361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3935 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 3994

Qy     421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3995 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 4054

Qy     481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4055 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 4114

Qy     541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4115 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 4174

Qy     601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4175 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 4234

Qy     661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4235 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 4294

Qy     721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4295 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 4354

Qy     781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4355 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 4414

Qy     841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4415 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 4474

Qy     901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4475 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 4534

Qy     961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4535 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCAG 4575


RESULT 8
US-10-095-718-3
; Sequence 3, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
;  PRIOR FILING DATE: 2001-08-22
;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3
;   LENGTH: 7914
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: rAAV vector with canine B-domain deleted factor
;   OTHER INFORMATION: VIII
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (435)...(4730)
US-10-095-718-3

  Query Match             84.1%;  Score 841.6;  DB 12;  Length 7914;
  Best Local Similarity   90.1%;  Pred. No. 2.7e-246;
  Matches  901;  Conservative    0;  Mismatches   99;  Indels    0;  Gaps    0;

Qy       1 ATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTC 60
          |||||| | |||||||||||||||||||||||||||||||| |||||||| ||||| ||||
Db    3541 ATCAAAAGGTTCGATGGTATCTGCTCAGCATGGGCAGCAACGAAAACATTCATTCCATTC 3600

Qy      61 ATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACA 120
          | |||||||||||||||||||||||||| |||||||||| |||||||||| | ||||
Db    3601 ACTTCAGTGGACATGTGTTCACTGTACGGAAAAAAGAGGAATATAAAATGGCAGTCTACA 3660

Qy     121 ATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGC 180
          | ||||||||||||||||||||||| |||||||| ||| |||||| ||||
Db    3661 ACCTCTATCCAGGTGTTTTTGAGACTGTGGAAATGCTACCATCCCAAGTTGGAATCTGGC 3720
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Qy    181 GGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGT 240
          || | ||||||||||| |||||||| || || || |||||||||| || ||||||||||
Db   3721 GGATAGAATGCCTTATCGGCGAGCACCTGCAAGCCGGGATGAGCACTCTGTTTCTGGTGT 3780

Qy    241 ACAGCAATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGA 300
          ||||||| |||||||||||||| ||||| |||||||| |||||||||||||||||||||
Db   3781 ACAGCAAGAAGTGTCAGACTCCACTGGGGATGGCTTCCGGACACATTAGAGATTTTCAGA 3840

Qy    301 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3841 TTACAGCTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCG 3900

Qy    361 GATCAATCAATGCCTGGAGCACCAAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGG 420
          |||||||||||||||||||||||||||| |||||||| |||||||||||||| ||||
Db   3901 GATCAATCAATGCCTGGAGCACCAAGGATCCCTTTTCCTGGATCAAGGTGGATCTCTTGG 3960

Qy    421 CACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCT 480
          |||| |||||||||||||||||| |||||||| ||||| |||||||||||||||||||
Db   3961 CACCGATGATTATTCACGGCATCATGACCCAGGGGGCCCGCCAGAAGTTCTCCAGCCTCT 4020

Qy    481 ACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAG 540
          || | |||||||||||||||||| ||||| ||||| || |||||||| | ||| ||||
Db   4021 ACGTGTCTCAGTTTATCATCATGTACAGTCTGGATGGCAACAAGTGGCACAGTTACCGAG 4080

Qy    541 GAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAAC 600
          | |||||||| || |||||||||||||||||||||||||| ||||||||||||||| ||||
Db   4081 GGAATTCCACGGGGACCTTAATGGTCTTCTTTGGCAACGTGGATTCATCTGGGATCAAAC 4140

Qy    601 ACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATA 660
          |||||||||||||||||| |||||||||  |||||||||||||||||||| ||||| |
Db   4141 ACAATATTTTTAACCCTCCGATTATTGCTCAGTACATCCGTTTGCACCCAACCCATTACA 4200

Qy    661 GCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGC 720
          |||| |||||||||||||||||||| |  |||||||||| || || ||||||||||||
Db   4201 GCATCCGCAGCACTCTTCGCATGGAGCTCTTGGGCTGTGACTTCAACAGTTGCAGCATGC 4260

Qy    721 CATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTA 780
          |  |||| |||||||||||||||||||||||||| ||||| ||||| || |||||| | |
Db   4261 CGCTGGGGATGGAGAGTAAAGCAATATCAGATGCTCAGATCACTGCCTCGTCCTACCTAA 4320

Qy    781 CCAATATGTTTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTA 840
          || |||| ||||||| |||||||||||  |||| || || |||| || || |||| ||
Db   4321 GCAGTATGCTTGCCACTTGGTCTCCTTCCCAAGCCCGGCTGCACCTGCAGGGCAGGACTA 4380

Qy    841 ATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGA 900
          ||||||||||||||||  ||||| |||||||||||||||||||||||||| |||||
Db   4381 ATGCCTGGAGACCTCAGGCAAATAACCCAAAAGAGTGGCTGCAAGTGGACTTCCGGAAGA 4440

Qy    901 CAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATG 960
          | |||||||||||||| |||| || ||||| || |||||||| || | ||||||||||
Db   4441 CCATGAAAGTCACAGGAATAACCACCCAGGGGGTGAAATCTCTCCTCATCAGCATGTATG 4500

Qy    961 TGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGGCCATCA 1000
          |||||||||||||||||||||| |||||||||||||||  |
Db   4501 TGAAGGAGTTCCTCATCTCCAGTAGTCAAGATGGCCATAA 4540
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RESULT 9
US-09-880-107-2275
; Sequence 2275, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
; APPLICANT: Vockley, Joseph G.
; APPLICANT: Scherf, Uwe
; APPLICANT: Gene Logic, Inc.
; TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
; FILE REFERENCE: 44921-5028-WO
; CURRENT APPLICATION NUMBER: US/09/880,107
; CURRENT FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/211,379
; PRIOR FILING DATE: 2000-06-14
; PRIOR APPLICATION NUMBER: US 60/237,054
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 3950
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2275
; LENGTH: 6909
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20020142981A1 M16967
US-09-880-107-2275

Query Match 19.8%; Score 198.2; DB 10; Length 6909;
Best Local Similarity 51.5%; Pred. No. 4.9e-50;
Matches 520; Conservative 0; Mismatches 468; Indels 21; Gaps 2;

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Qy      7 GGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCA 66
          || |  | |   | ||||| | ||| ||| ||    || |||| ||     |||| ||
Db   5564 GGGTGAGGTTACACCTGCTGAACATAGGCGGCTCCCAAGACATTCACGTGGTTCACTTTC 5623

Qy     67 GTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCT 126
            || ||    ||   | |      || |  || | |  | |  | |    ||
Db   5624 ACGGCCAGACCTTGCTGGAAAATGGCAATAAACAGCACCAGTTAGGGGTCTGGCCCCTTC 5683

Qy    127 ATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGG 186
            || |||   ||| | ||  | ||||||    |||| ||| ||||    ||||   |
Db   5684 TGCCTGGTTCATTTAAAACTCTTGAAATGAAGGCATCAAAACCTGGCTGGTGGCTCCTAA 5743

Qy    187 AATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCA 246
          |        |||| ||  | |     || ||||||   || |  |||||  |       ||
Db   5744 ACACAGAGGTTGGAGAAAACCAGAGAGCAGGGATGCAAACGCCATTTCTTATCATGGACA 5803

Qy    247 ATAAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAG 306
            | |||  ||  ||  |||||  |     ||||   |||    ||||  |||||  |  |
Db   5804 GAGACTGTAGGATGCCAATGGGACTAAGCACTGGTATCATATCTGATTCACAGATCAAGG 5863

Qy    307 CTTCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAA 366
          ||||||      ||  | ||||  || |   | || ||| |  |  ||  |||||
Db   5864 CTTCAGAGTTTCTGGGTTACTGGGAGCCCAGATTAGCAAGATTAAACAATGGTGGATCTT 5923

Qy    367 TCAATGCCTGGAGCACCAAGGAGC----------------CCTTTTCTTGGATCAAGG 408
            |||||  |||||     |  | |                |     |||||||| |||
Db   5924 ATAATGCTTGGAGTGTAGAAAAACTTGCAGCAGAATTTGCCTCTAAACCTTGGATCCAGG 5983

Qy    409 TGGATCTGTTGGCACCAATGATTATTCACGGCATCAAGACCCAGGGTGCCCGTCAGAAGT 468
          ||||  ||        | | || ||     || ||| ||||||| ||||||   ||  |
Db   5984 TGGACATGCAAAAGGAAGTCATAATCACAGGGATCCAGACCCAAGGTGCCAAACACTACC 6043

Qy    469 TCTCCAGCCTCTACATCTCTCAGTTTATCATCATGTATAGTCTTGATGGGAAGAAGTGGC 528
          |       |  ||| | | |  ||||      |    || |||    |   ||  || ||||
Db   6044 TGAAGTCCTGCTATACCACAGAGTTCTATGTAGCTTACAGTTCCAACCAGATCAACTGGC 6103

Qy    529 AGACTTATCGAGGAAATTCCACTGGAACCTTAATGGTCTTCTTTGGCAATGTGGATTCAT 588
          |||  |    ||| ||   |||  | |   | |||   ||   |||||||   ||| | |
Db   6104 AGATCTTCAAAGGGAACAGCACAAGGAATGTGATGTATTTTAATGGCAATTCAGATGCCT 6163

Qy    589 CTGGGATAAAACACAATATTTTTAACCCTCCAATTATTGCTCGATACATCCGTTTGCACC 648
          ||   |||||| | |||   ||| |||| || ||| | ||| |||| ||  |  |   |
Db   6164 CTACAATAAAAGAGAATCAGTTTGACCCACCTATTGTGGCTAGATATATTAGGATCTCTC 6223

Qy    649 CAACTCATTATAGCATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATA 708
          ||||||         |   |    || |||||  ||||  ||   || |||||  |||||
Db   6224 CAACTCGAGCCTATAACAGACCTACCCTTCGATTGGAACTGCAAGGTTGTGAGGTAAATG 6283

Qy    709 GTTGCAGCATGCCATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTT 768
          | ||   ||  ||  |||| ||||| | |  |   |||  | |    || || || ||||
Db   6284 GATGTTCCACACCCCTGGGTATGGAAAATGGAAAGATAGAAAACAAGCAAATCACAGCTT 6343

Qy    769 CATCCTACTTTACCAATATGT---TTGCCACCTGGTCTCCTTCAAAAGCTCGACTTCACC 825
          | || |     |    | ||     |   ||||   || |     || || ||  |
Db   6344 CTTCGTTTAAGAAATCTTGGTGGGGAGATTACTGGGAACCCTTCCGTGCCCGTCTGAATG 6403

Qy    826 TCCAAGGGAGGAGTAATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAG 885
            ||| ||  |    |||||||||  | |  |||  || ||    ||  |||||||  ||
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Db    6404 CCCAGGGACGTGTGAATGCCTGGCAAGCCAAGGCAAACAACAATAAGCAGTGGCTAGAAA 6463

Qy     886 TGGACTTCCAGAAGACAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGC 945
           | ||  | |  ||||  | |||  | || | | | |  || |||||    || ||||||
Db    6464 TTGATCTACTCAAGATCAAGAAGATAACGGCAATTATAACACAGGGCTGCAAGTCTCTGT 6523

Qy     946 TTACCAGCATGTATGTGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGG 994
             |    |||||||| |||   |    ||||  |  |||| |  | ||
Db    6524 CCTCTGAAATGTATGTAAAGAGCTATACCATCCACTACAGTGAGCAGGG 6572


RESULT 10
US-09-974-298-167
; Sequence 167, Application US/09974298
; Patent No. US20020156263A1
; GENERAL INFORMATION:
;  APPLICANT: Chen, Huei-Mei
;  TITLE OF INVENTION: GENES EXPRESSED IN BREAST CANCER
;  FILE REFERENCE: PA-0037 P
;  CURRENT APPLICATION NUMBER: US/09/974,298
;  CURRENT FILING DATE:  2001-10-04
;  PRIOR APPLICATION NUMBER: 60/238,331
;  PRIOR FILING DATE: 2000-05-10
;  NUMBER OF SEQ ID NOS: 194
;  SOFTWARE: PERL Program
; SEQ ID NO 167
;   LENGTH: 4599
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   OTHER INFORMATION: Incyte ID No. US20020156263A1 235636.1
US-09-974-298-167

  Query Match             9.2%;  Score 92.4;  DB 9;  Length 4599;
  Best Local Similarity   48.1%;  Pred. No. 7.6e-18;
  Matches  301;  Conservative    0;  Mismatches  316;  Indels    9;  Gaps    1;

Qy     384 AAGGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAATGATTATTCACGGCATC 443
           || ||    |   | |||||  || |  || ||        |||||   ||   ||  |
Db     941 AATGACAGATGGCCGTGGATTCAGATAAATTTGCAAAGGAAAATGAGAGTTACTGGTGTG 1000

Qy     444 AAGACCCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCTACATCTCTCAGTTTATCATCATG 503
           |  ||||| || |||     ||       |      || ||       |  |  ||
Db    1001 ATTACCCAAGGAGCCAAGAGGATTGGAAGCCCAGAGTATATAAAATCCTACAAAATTGCC 1060

Qy     504 TATAGTCTTGATGGGAAGAAGTGGCAGACTTATCGAGGAAATTCCACTGGAACCTTAATG 563
           || |||  |||||| ||||  |||   |  ||   ||  ||   |||          |||
Db    1061 TACAGTAATGATGGAAAGACTTGGGCAATGTACAAAGTGAAAGGCACCAATGAAGACATG 1120

Qy     564 GTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAACACAATATTTTTAACCCTCCAATT 623
           || ||   ||| ||  | |||          | |    ||   ||| |  || || ||
Db    1121 GTGTTTCGTGGAAACATTGATAACAACACTCCATATGCTAACTCTTTCACACCCCCCATA 1180

Qy     624 ATTGCTCGATACATCCGTTTGCACCCAACTCATTATAGCATTCGCAGCACTCTTCGCATG 683
           |  ||||  ||  |  |  |  | ||      || | | |  |   |||||  | || |||
Db    1181 AAAGCTCAGTATGTAAGACTCTATCCCCAAGTTTGTCGAAGACATTGCACTTTGCGAATG 1240

Qy     684 GAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGCCATTGGGAATGGAGAGTAAAGCA 743
           ||  |  | ||||||||  |     ||||      |||  |||| ||| |       |
Db    1241 GAACTTCTTGGCTGTGAACTGTCGGGTTGTTCTGAGCCTCTGGGTATGAAATCAGGACAT 1300

Qy     744 ATATCAGATGCACAGATTACTGCTTCATCCTACT---------TTACCAATATGTTTGCC 794
           |||  |||    ||||| ||||| ||  |  ||        | | ||      |   |
Db    1301 ATACAAGACTATCAGATCACTGCCTCCAGCATCTTCAGAACGCTCAACATGGACATGTTC 1360

Qy     795 ACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTAATGCCTGGAGACCT 854
           || |||   ||    |||||||| ||  ||   ||||| |     ||||||||||   ||
Db    1361 ACTTGGGAACCAAGGAAAGCTCGGCTGGACAAGCAAGGCAAAGTGAATGCCTGGACCTCT 1420

Qy     855 CAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGACAATGAAAGTCACA 914
             ||| | |    | | ||| | || |||||  | |      |||  ||||| ||
Db    1421 GGCCACAATGACCAGTCACAATGGTTACAGGTGGATCTTCTTGTTCCAACCAAAGTGACT 1480

Qy     915 GGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGTATGTGAAGGAGTTCCTC 974
           ||  | | ||| || ||||  |||  | |   |         || |||       | |
Db    1481 GGCATCATTACACAAGGAGCTAAAGATTTTGGTCATGTACAGTTTGTTGGCTCCTACAAA 1540

Qy     975 ATCTCCAGCAGTCAAGATGGCCATCA 1000
             |  |  |||  | |||||  | ||
Db    1541 CTGGCTTACAGCAATGATGGAGAACA 1566


RESULT 11
US-10-003-132-12
; Sequence 12, Application US/10003132
; Publication No. US20020192750A1
; GENERAL INFORMATION:
;  APPLICANT: Fox, Brian A.
;  APPLICANT:  Gao, Zeren
;  APPLICANT:  Shoemaker, Kimberly E.
;  TITLE OF INVENTION: NEUROPILIN HOMOLOG ZCUB5
;  FILE REFERENCE: 00-62
;  CURRENT APPLICATION NUMBER: US/10/003,132
;  CURRENT FILING DATE:  2001-11-15
;  PRIOR APPLICATION NUMBER: US 60/249,004
;  PRIOR FILING DATE: 2000-11-15
;  NUMBER OF SEQ ID NOS: 19
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 12
;   LENGTH: 2145
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: degenerate nucleotide sequence
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(2145)
;   OTHER INFORMATION: n = A,T,C or G
US-10-003-132-12

  Query Match             7.6%;  Score 76;  DB 9;  Length 2145;
  Best Local Similarity   32.4%;  Pred. No. 5.1e-13;
  Matches  122;  Conservative   64;  Mismatches  191;  Indels    0;  Gaps    0;

Qy     329 GGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAAGGA 388
           :|| :     | :| :  :   |  :  |  |  ::     |: : :: | :| : |  |
Db     867 RGCNMGNYTNCARGAYCARGGNCCNWSNTGGGCNWSNGGNGAYWSNWSNAAYAAYCAYAA 926

Qy     389 GCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAATGATTATTCACGGCATCAAGAC 448
           :||       ||| |  |: | ||::|    | :  :| :|  ||:   || ||::  ||
Db     927 RCCNMGNGARTGGYTNGARATHGAYYTNGGNGARAARAARAARATHACNGGNATHMGNAC 986

Qy     449 CCAGGGTGCCCGTCAGAAGTTCTCCAGCCTCTACATCTCTCAGTTTATCATCATGTATAG 508
             ||  :    ||::    :| :| : |:||: |       ||: | || |  | :|
Db     987 NACNGGNWSNACNCARWSNAAYTTYAAYTTYTAYGTNAARWSNTTYGTNATGAAYTTYAA 1046

Qy     509 TCTTGATGGGAAGAAGTGGCAGACTTATCGAGGAAATTCCACTGGAACCTTAATGGTCTT 568
             : |:   :  ||:||| |:|| ||:  :|| | :   | :        :| :|| ||
Db    1047 RAAYAAYAAYWSNAARTGGAARACNTAYAARGGNATHGTNAAYAAYGARGARAARGTNTT 1106

Qy     569 CTTTGGCAATGTGGATTCATCTGGGATAAAACACAATATTTTTAACCCTCCAATTATTGC 628
           :   || ||:    |:|     |        || ||:| :||:| :|| || ||: | ||
Db    1107 YCARGGNAAYWSNAAYTTYMGNGAYCCNGTNCARAAYAAYTTYATHCCNCCNATHGTNGC 1166

Qy     629 TCGATACATCCGTTTGCACCCAACTCATTATAGCATTCGCAGCACTCTTCGCATGGAGTT 688
           :| ||: | :|  |    ||      |    :  :| | : | :|     | ||::|
Db    1167 NMGNTAYGTNMGNGTNGTNCCNCARACNTGGCAYCARMGNATHGCNYTNAARGTNGARYT 1226

Qy     689 GATGGGCTGTGATTTAA 705
           || || ||: |  |:|
Db    1227 NATHGGNTGYCARATHA 1243


RESULT 12
US-09-880-107-3020
; Sequence 3020, Application US/09880107
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; Patent No. US20020142981A1
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,107
;  CURRENT FILING DATE:  2001-06-14
;  PRIOR APPLICATION NUMBER: US 60/211,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3020
;   LENGTH: 1270
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020142981A1 S56151
US-09-880-107-3020

  Query Match             7.2%;  Score 72.4;  DB 10;  Length 1270;
  Best Local Similarity   51.6%;  Pred. No. 4.8e-12;
  Matches  197;  Conservative    0;  Mismatches  176;  Indels    9;  Gaps    1;

Qy    561 ATGGTCTTCTTTGGCAATGTGGATTCATCTGGGATAAAACACAATATTTTTAACCCTCCA 620
          | ||  ||  | || ||   | |   |   | | |  |   |||  | ||| |  | ||
Db     31 AAGGAGTTTGTGGGTAACTGGAACAAAAACGCGGTGCATGTCAACCTGTTTGAGACCCCT 90

Qy    621 ATTATTGCTCGATACATCCGTTTGCACCCAACTCATTATAGCATTCGCAGCACTCTTCGC 680
            |    ||||  ||| |  | ||| |||| ||    |    ||    | ||||||| |||
Db     91 GTGGAGGCTCAGTACGTGAGATTGTACCCCACGAGCTGCCACACGGCCTGCACTCTGCGC 150

Qy    681 ATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGCCATTGGGAATGGAGAGTAAA 740
          | ||| |  ||||||||||  | ||  | |||  ||  ||  ||||  || ||| |||
Db    151 TTTGAGCTACTGGGCTGTGAGCTGAACGGATGCGCCAATCCCCTGGGCCTGAAGAATAAC 210

Qy    741 GCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTACCAATATG---------TTT 791
             ||  | ||    ||||| || || ||   |  ||  |  |    |         |
Db    211 AGCATCCCTGACAAGCAGATCACGGCCTCCAGCAGCTACAAGACCTGGGGCTTGCATCTC 270

Qy    792 GCCACCTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGAGTAATGCCTGGAGA 851
            || ||||   || ||  | || || ||  ||   || || |     || ||||||
Db    271 TTCAGCTGGAACCCCTCCTATGCACGGCTGGACAAGCAGGGCAACTTCAACGCCTGGGTT 330

Qy    852 CCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGAAGACAATGAAAGTC 911
          |   |    |   |    |  |||||||||| ||||||  |         | | | | ||
Db    331 GCGGGGAGCTACGGTAACGATCAGTGGCTGCAGGTGGACCTGGGCTCCTCGAAGGAGGTG 390

Qy    912 ACAGGAGTAACTACTCAGGGAG 933
          |||||  | |  || ||||| |
Db    391 ACAGGCATCATCACCCAGGGGG 412


RESULT 13
US-09-960-352-13850
; Sequence 13850, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 13850
;   LENGTH: 357
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   OTHER INFORMATION: Clone ID: 59-LIB34-008-Q1-E1-G12
US-09-960-352-13850

  Query Match             6.9%;  Score 69.2;  DB 10;  Length 357;
  Best Local Similarity   54.8%;  Pred. No. 2.3e-11;
  Matches  137;  Conservative    0;  Mismatches  113;  Indels    0;  Gaps    0;

Qy    129 CCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAA 188
          || |||   ||| | ||  | ||||||    |||| ||| ||||    ||||   | ||
Db     34 CCTGGTTCATTTAAAACTCTTGAAATGAAGGCATCAAAACCTGGCTGGTGGCTCCTAGAC 93

Qy    189 TGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAAT 248
                |||| ||   ||     || ||||||  ||||  |||||  |     ||
Db     94 ACGGAAGTTGGAGAAATTCAGAGAGCAGGGATGCAGACACCATTTCTCATTGTAGACAGA 153

Qy    249 AAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCT 308
            | ||| |||  ||  ||||| |     |||| |  ||    || |  |||||   |||
Db    154 GAATGTAAGATGCCAATGGGACTAAGCACTGGCCTGATAGCTGACTCACAGATCCAGGCT 213

Qy    309 TCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATC 368
          || |     |  ||  | ||||  || ||  | || ||  |  |  ||   ||||||  |
Db    214 TCTGAGTTTTGGGGTTATTGGGAACCCAAATTAGCAAGGTTAAACAATGGTGGATCATAC 273

Qy    369 AATGCCTGGA 378
          |||||  ||||
Db    274 AATGCTTGGA 283


RESULT 14
US-09-960-352-8113
; Sequence 8113, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 8113
;   LENGTH: 389
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   OTHER INFORMATION: Clone ID: 35-LIB2809-021-Q1-E1-A4
US-09-960-352-8113

  Query Match             6.1%;  Score 61.2;  DB 10;  Length 389;
  Best Local Similarity   51.2%;  Pred. No. 6.8e-09;
  Matches  175;  Conservative    0;  Mismatches  158;  Indels    9;  Gaps    1;

Qy    662 CATTCGCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGCC 721
          ||   || |||| || ||| | ||  |  | ||||||| || ||| | ||||     ||
Db     31 CAAGGGCTGCACCCTCCGCTTTGAACTCCTTGGCTGTGAGTTGAATGGATGCACTGAACC 90

Qy    722 ATTGGGAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTAC 781
            | ||  || ||  |||  | ||  |  |    ||||| || || ||   |||||  |
Db     91 CCTAGGCCTGAAGGATAATACCATCCCCAACAAGCAGATCACAGCCTCCAGCTACTACAA 150

Qy    782 CAATATGTTTGCCAC---------CTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGG 832
          |   |    |           ||||| ||| |   | || ||||  |    || ||
Db    151 AACCTGGGGCCTGAGTGCCTTTAGCTGGTTTCCCTACTACGCACGACTGGATAATCAGGG 210

Qy    833 GAGGAGTAATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTT 892
          | |   || |||||||   | |||   || | | |    ||||||||||  | ||| |
Db    211 CAAGTTCAACGCCTGGACCGCCCAGACCAACAGTGCCTCTGAGTGGCTGCAGATTGACCT 270
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Qy    893 CCAGAAGACAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAG 952
                  | |  |||||| ||  | |  || ||||| |    |    |      ||
Db    271 GGGCTCCCAGAAGCGAGTCACGGGCATCATCACCCAGGGTGCCCGAGACTTTGGCCACAT 330

Qy    953 CATGTATGTGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGG 994
              ||||||   |  | |    |  ||    || | |||||
Db    331 TCAATATGTGGCTGCCTACAGGGTGGCCTATGGTGATGATGG 372


RESULT 15
US-09-960-352-3109
; Sequence 3109, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 3109
;   LENGTH: 367
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   OTHER INFORMATION: Clone ID: 14-LIB2809-023-Q1-E1-D5
US-09-960-352-3109

  Query Match             6.1%;  Score 61;  DB 10;  Length 367;
  Best Local Similarity   51.3%;  Pred. No. 7.6e-09;
  Matches  173;  Conservative    0;  Mismatches  155;  Indels    9;  Gaps    1;

Qy    667 GCAGCACTCTTCGCATGGAGTTGATGGGCTGTGATTTAAATAGTTGCAGCATGCCATTGG 726
          || |||| || ||| | ||  |  | |||||||| || ||| | ||||     ||  | |
Db     30 GCTGCACCCTCCGCTTTGAACTCCTTGGCTGTGAGTTGAATGGATGCACTGAACCCCTAG 89

Qy    727 GAATGGAGAGTAAAGCAATATCAGATGCACAGATTACTGCTTCATCCTACTTTACCAATA 786
          |  || ||  |||  | ||  |  |    ||||| || || ||   |||||  |  |
Db     90 GCCTGAAGGATAATACCATCCCCAACAAGCAGATCACAGCCTCCAGCTACTACAAAACCT 149

Qy    787 TGTTTGCCAC---------CTGGTCTCCTTCAAAAGCTCGACTTCACCTCCAAGGGAGGA 837
           |       |          ||||| ||| |   | || |||||  |    || || | |
Db    150 GGGGCCTGAGTGCCTTTAGCTGGTTTCCCTACTACGCACGACTGGATAATCAGGGCAAGT 209

Qy    838 GTAATGCCTGGAGACCTCAGGTGAATAATCCAAAAGAGTGGCTGCAAGTGGACTTCCAGA 897
            || |||||||   | |||   || | | |    ||||||||||  | ||| |
Db    210 TCAACGCCTGGACCGCCCAGACCAACAGTGCCTCTGAGTGGCTGCAGATTGACCTGGGCT 269

Qy    898 AGACAATGAAAGTCACAGGAGTAACTACTCAGGGAGTAAAATCTCTGCTTACCAGCATGT 957
          | |  |||||| ||  | |  || ||||| |    |    |      ||     |
Db    270 CCCAGAAGCGAGTCACGGGCATCATCACCCAGGGTGCCCGAGACTTTGGCCACATTCAAT 329

Qy    958 ATGTGAAGGAGTTCCTCATCTCCAGCAGTCAAGATGG 994
          |||||   |  | |    |  ||    || | |||||
Db    330 ATGTGGCTGCCTACAGGGTGGCCTATGGTGATGATGG 366


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Job time : 144 secs
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GenCore version 5.1.3

OM protein - protein search, using sw model

Run on: January 5, 2003, 04:48:07 ; Search time 24 Seconds
(without alignments)
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Title: US-09-740-211-15
Perfect score: 75
Sequence: 1 SFSQNPPVLKRHQR 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	75	100.0	14	2	US-08-634-001C-1	Sequence 1, Appli
2	75	100.0	14	4	US-09-470-618-15	Sequence 15, Appl
3	75	100.0	1438	4	US-09-209-916-1	Sequence 1, Appli
4	65	86.7	1661	2	US-08-882-083-2	Sequence 2, Appli
5	65	86.7	1661	2	US-08-558-107-2	Sequence 2, Appli
6	65	86.7	1661	4	US-09-243-539-2	Sequence 2, Appli
7	65	86.7	2332	1	US-07-864-004B-4	Sequence 4, Appli
8	65	86.7	2332	1	US-08-251-937A-4	Sequence 4, Appli
9	65	86.7	2332	1	US-08-212-133A-2	Sequence 2, Appli
10	65	86.7	2332	1	US-08-276-594A-2	Sequence 2, Appli
11	65	86.7	2332	1	US-08-474-503-2	Sequence 2, Appli
12	65	86.7	2332	2	US-08-670-707A-2	Sequence 2, Appli
13	65	86.7	2332	4	US-09-037-601-2	Sequence 2, Appli
14	65	86.7	2332	4	US-09-324-867-3	Sequence 3, Appli
15	65	86.7	2332	4	US-09-315-179-2	Sequence 2, Appli
16	65	86.7	2332	4	US-09-523-656-2	Sequence 2, Appli
17	65	86.7	2332	5	PCT-US93-03275-4	Sequence 4, Appli
18	65	86.7	2332	5	PCT-US94-13200-2	Sequence 2, Appli
19	65	86.7	2351	1	US-08-121-202-2	Sequence 2, Appli
20	65	86.7	2351	1	US-08-366-851A-2	Sequence 2, Appli
21	65	86.7	2351	6	5171844-2	Patent No. 5171844
22	65	86.7	2351	6	5422260-1	Patent No. 5422260
23	58	77.3	1471	1	US-08-683-839B-3	Sequence 3, Appli
24	56	74.7	2304	4	US-09-324-867-4	Sequence 4, Appli
25	56	74.7	2319	1	US-08-212-133A-8	Sequence 8, Appli
26	56	74.7	2319	1	US-08-474-503-6	Sequence 6, Appli
27	56	74.7	2319	2	US-08-670-707A-6	Sequence 6, Appli
28	56	74.7	2319	4	US-09-037-601-6	Sequence 6, Appli
29	56	74.7	2319	4	US-09-315-179-6	Sequence 6, Appli
30	56	74.7	2319	4	US-09-523-656-28	Sequence 28, Appl
31	56	74.7	2319	5	PCT-US94-13200-6	Sequence 6, Appli
32	55	73.3	34	3	US-08-441-935-30	Sequence 30, Appl
33	55	73.3	34	4	US-08-441-943-30	Sequence 30, Appl
34	54	72.0	24	4	US-09-523-656-32	Sequence 32, Appl
35	54	72.0	1467	4	US-09-523-656-38	Sequence 38, Appl
36	54	72.0	2343	4	US-09-324-867-2	Sequence 2, Appli
37	50	66.7	9	4	US-08-441-943-34	Sequence 34, Appl
38	47	62.7	868	1	US-07-864-004B-6	Sequence 6, Appli
39	47	62.7	868	1	US-08-251-937A-6	Sequence 6, Appli
40	47	62.7	868	1	US-08-212-133A-3	Sequence 3, Appli
41	47	62.7	1090	5	PCT-US93-03275-6	Sequence 6, Appli
42	47	62.7	2115	4	US-09-324-867-5	Sequence 5, Appli
43	47	62.7	2133	2	US-08-670-707A-37	Sequence 37, Appl
44	47	62.7	2133	4	US-09-037-601-37	Sequence 37, Appl
45	47	62.7	2133	4	US-09-315-179-37	Sequence 37, Appl

ALIGNMENTS

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RESULT 1
US-08-634-001C-1
; Sequence 1, Application US/08634001C
; Patent No. 5952198
;  GENERAL INFORMATION:
;    APPLICANT:  Chan, Sham-Yuen
;    TITLE OF INVENTION:  Production of Recombinant Factor VIII
;    TITLE OF INVENTION:  in the Presence of Liposome-like
;    TITLE OF INVENTION:  Substances of Mixed Composition
;    NUMBER OF SEQUENCES:  1
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Bayer Corporation
;      STREET:  800 Dwight Way
;      STREET:  P. O. Box 1986
;      CITY:  Berkeley
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94701-1986
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette, 3.50 inch, 1.44Mb Storage
;      COMPUTER:  IBM
;      OPERATING SYSTEM:  DOS
;      SOFTWARE:  WordPerfect 6.1
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/634,001C
;      FILING DATE:
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/434,900
;      FILING DATE:  May 4, 1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Giblin, James A.
;      REGISTRATION NUMBER:  25772
;      REFERENCE/DOCKET NUMBER:  MSB-7226CIP
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (510)705-7910
;      TELEFAX:  (510)705-7904
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  14
;      TYPE:  amino acid
;      STRANDEDNESS:  single strand
;      TOPOLOGY:  linear
;    MOLECULE TYPE:
;      DESCRIPTION:  peptide
US-08-634-001C-1

  Query Match             100.0%;  Score 75;  DB 2;  Length 14;
  Best Local Similarity   100.0%;  Pred. No. 4.7e-07;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;
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Qy      1 SFSQNPPVLKRHQR 14
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Db      1 SFSQNPPVLKRHQR 14


RESULT 2
US-09-470-618-15
; Sequence 15, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION: by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/470,618
;  CURRENT FILING DATE: 1999-12-22
;  EARLIER APPLICATION NUMBER: 09/364,862
;  EARLIER FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 15
;   LENGTH: 14
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-470-618-15

  Query Match             100.0%;  Score 75;  DB 4;  Length 14;
  Best Local Similarity   100.0%;  Pred. No. 4.7e-07;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 SFSQNPPVLKRHQR 14
          ||||||||||||||
Db      1 SFSQNPPVLKRHQR 14


RESULT 3
US-09-209-916-1
; Sequence 1, Application US/09209916
; Patent No. 6358703
; GENERAL INFORMATION:
;  APPLICANT: Cho, Myung-Sam
;  APPLICANT: Chan, Sham-Yuen
;  APPLICANT: Kelsey, William
;  APPLICANT: Yee, Helena
;  TITLE OF INVENTION: Expression System for Factor VIII
;  FILE REFERENCE: MSB-7255
;  CURRENT APPLICATION NUMBER: US/09/209,916
;  CURRENT FILING DATE: 1998-12-10
;  NUMBER OF SEQ ID NOS: 2
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 1438
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Derived from
;   OTHER INFORMATION: human factor VIII sequence
US-09-209-916-1

  Query Match             100.0%;  Score 75;  DB 4;  Length 1438;
  Best Local Similarity   100.0%;  Pred. No. 6.9e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 SFSQNPPVLKRHQR 14
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Db    741 SFSQNPPVLKRHQR 754


RESULT 4
US-08-882-083-2
; Sequence 2, Application US/08882083
; Patent No. 5869292
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/882,083
;      FILING DATE:
;      CLASSIFICATION:  514
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1661 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-08-882-083-2

  Query Match             86.7%;  Score 65;  DB 2;  Length 1661;
  Best Local Similarity   100.0%;  Pred. No. 0.0045;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      3 SQNPPVLKRHQR 14
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Db    966 SQNPPVLKRHQR 977


RESULT 5
US-08-558-107-2
; Sequence 2, Application US/08558107
; Patent No. 5910481
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
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;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/558,107
;      FILING DATE:  13-NOV-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1661 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-08-558-107-2

  Query Match             86.7%;  Score 65;  DB 2;  Length 1661;
  Best Local Similarity   100.0%;  Pred. No. 0.0045;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db     966 SQNPPVLKRHQR 977


RESULT 6
US-09-243-539-2
; Sequence 2, Application US/09243539
; Patent No. 6130203
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/243,539
;      FILING DATE:
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1661 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-09-243-539-2

  Query Match             86.7%;  Score 65;  DB 4;  Length 1661;
  Best Local Similarity   100.0%;  Pred. No. 0.0045;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db     966 SQNPPVLKRHQR 977


RESULT 7
US-07-864-004B-4
; Sequence 4, Application US/07864004B
; Patent No. 5364771
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  6
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/864,004B
;      FILING DATE:  07 APRIL 1992
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU106
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6508
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver cDNA sequence
US-07-864-004B-4

  Query Match             86.7%;  Score 65;  DB 1;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
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Db    1637 SQNPPVLKRHQR 1648


RESULT 8
US-08-251-937A-4
; Sequence 4, Application US/08251937A
; Patent No. 5583209
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
```

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;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  10
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/251,937A
;      FILING DATE:  31-MAY-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106DIV
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6367
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver cDNA sequence
US-08-251-937A-4

  Query Match             86.7%;  Score 65;  DB 1;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db    1637 SQNPPVLKRHQR 1648


RESULT 9
US-08-212-133A-2
; Sequence 2, Application US/08212133A
; Patent No. 5663060
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/212,133A
;      FILING DATE:  March 11, 1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU/76677
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-572-6508
;      TELEFAX:  404-572-6555
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  YES
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver cDNA sequence
US-08-212-133A-2

  Query Match             86.7%;  Score 65;  DB 1;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db    1637 SQNPPVLKRHQR 1648


RESULT 10
US-08-276-594A-2
; Sequence 2, Application US/08276594A
; Patent No. 5693499
;  GENERAL INFORMATION:
;    APPLICANT:  YONEMURA, Hiroshi
;    APPLICANT:  TAJIMA, Yoshitaka
;    APPLICANT:  SUGAWARA, Keishin
;    APPLICANT:  MASUDA, Kenichi
;    TITLE OF INVENTION:  PROCESS FOR PREPARING HUMAN COAGULATION
;    TITLE OF INVENTION:  FACTOR VIII PROTEIN COMPLEX
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/276,594A
;      FILING DATE:  18-JUL-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/950,191
;      FILING DATE:  24-SEP-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 243262/1991
;      FILING DATE:  24-SEP-1991
;    ATTORNEY/AGENT INFORMATION:
```

```
;      NAME:  WEGNER, Harold C.
;      REGISTRATION NUMBER:  25,258
;      REFERENCE/DOCKET NUMBER:  74129/195/AOPA
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-08-276-594A-2

  Query Match             86.7%;  Score 65;  DB 1;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db    1637 SQNPPVLKRHQR 1648


RESULT 11
US-08-474-503-2
; Sequence 2, Application US/08474503
; Patent No. 5744446
;  GENERAL INFORMATION:
;    APPLICANT:  Emory University
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street, Suite 2800
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/474,503
;      FILING DATE:  07-JUN-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106CIP(3)
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6500
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  YES
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver cDNA sequence
US-08-474-503-2

  Query Match             86.7%;  Score 65;  DB 1;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
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Db    1637 SQNPPVLKRHQR 1648


RESULT 12
US-08-670-707A-2
; Sequence 2, Application US/08670707A
; Patent No. 5859204
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/670,707A
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Greenlee, Lorance L.
;      REGISTRATION NUMBER:  27,894
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  YES
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapiens
;      TISSUE TYPE:  Liver
US-08-670-707A-2

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  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db    1637 SQNPPVLKRHQR 1648


RESULT 13
US-09-037-601-2
; Sequence 2, Application US/09037601
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; Patent No. 6180371
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  40
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;      STREET:  5370 Manhattan Circle Suite 201
;      CITY:  Boulder
;      STATE:  Colorado
;      COUNTRY:  USA
;      ZIP:  80303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/037,601
;      FILING DATE:  26-JUN-1996
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US94/13200
;      FILING DATE:  15-NOV-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/212,133
;      FILING DATE:  11-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Ferber, Donna M.
;      REGISTRATION NUMBER:  33,878
;      REFERENCE/DOCKET NUMBER:  75-95F
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  303/499-8080
;      TELEFAX:  303/499-8089
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2332 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  not relevant
;    MOLECULE TYPE:  protein
;    HYPOTHETICAL:  YES
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapiens
;      TISSUE TYPE:  Liver
US-09-037-601-2

  Query Match             86.7%;  Score 65;  DB 4;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0066;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
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Db    1637 SQNPPVLKRHQR 1648


RESULT 14
US-09-324-867-3
; Sequence 3, Application US/09324867A
; Patent No. 6251632
; GENERAL INFORMATION:
;  APPLICANT: Lillicrap, David
;  APPLICANT: Cameron, Cherie
;  APPLICANT: No. 6251632ley, Colleen
;  APPLICANT: Horrocks, L. Suzanne Hoyle
;  APPLICANT: Hough, Christine
;  TITLE OF INVENTION: Canine Factor VIII Gene, Protein and Methods of Use
;  FILE REFERENCE: 1669.0010002/JAG/BJD
;  CURRENT APPLICATION NUMBER: US/09/324,867A
;  CURRENT FILING DATE: 1999-06-03
;  EARLIER APPLICATION NUMBER: 09/035,141
;  EARLIER FILING DATE: 1998-03-059
;  EARLIER APPLICATION NUMBER: 60/039,953
;  EARLIER FILING DATE: 1997-03-06
;  NUMBER OF SEQ ID NOS: 63
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 3
;   LENGTH: 2332
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-324-867-3

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Qy       3 SQNPPVLKRHQR 14
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Db    1637 SQNPPVLKRHQR 1648


RESULT 15
US-09-315-179-2
; Sequence 2, Application US/09315179
; Patent No. 6376463
; GENERAL INFORMATION:
;  APPLICANT: Lollar, John S
;  TITLE OF INVENTION: Modified Factor VIII
;  FILE REFERENCE: 75-95H
;  CURRENT APPLICATION NUMBER: US/09/315,179
;  CURRENT FILING DATE: 1999-05-20
;  EARLIER APPLICATION NUMBER: U.S. 09/037,601
;  EARLIER FILING DATE: 1998-03-10
;  EARLIER APPLICATION NUMBER: U.S. 08/670,707
;  EARLIER FILING DATE: 1996-06-26
;  EARLIER APPLICATION NUMBER: PCT/US97/11155
;  EARLIER FILING DATE: 1997-06-26
;  EARLIER APPLICATION NUMBER: PCT/US94/13200
;  EARLIER FILING DATE: 1994-11-15
;  EARLIER APPLICATION NUMBER: U.S. 08/212,133
;  EARLIER FILING DATE: 1994-03-11
;  EARLIER APPLICATION NUMBER: U.S. 07/864,004
;  EARLIER FILING DATE: 1992-04-07
;  NUMBER OF SEQ ID NOS: 40
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
;   LENGTH: 2332
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-315-179-2

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  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       3 SQNPPVLKRHQR 14
           ||||||||||||
Db    1637 SQNPPVLKRHQR 1648


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Job time : 26 secs
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GenCore version 5.1.3

OM protein - protein search, using sw model

Run on: January 5, 2003, 06:13:09 ; Search time 17 Seconds
(without alignments)
15.607 Million cell updates/sec

Title: US-09-740-211-15
Perfect score: 75
Sequence: 1 SFSQNPPVLKRHQR 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 117078 seqs, 18951520 residues

Total number of hits satisfying chosen parameters: 117078

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA:*
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14: /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	75	100.0	14	9	US-10-007-968-15	Sequence 15, Appl
2	75	100.0	14	10	US-09-740-211-15	Sequence 15, Appl
3	75	100.0	1438	12	US-10-006-091-1	Sequence 1, Appli
4	75	100.0	1438	12	US-10-047-257-1	Sequence 1, Appli
5	65	86.7	2332	9	US-09-957-641-2	Sequence 2, Appli
6	58	77.3	1471	12	US-10-095-718-2	Sequence 2, Appli
7	55	73.3	34	10	US-09-748-062-30	Sequence 30, Appl
8	50	66.7	9	10	US-09-748-062-34	Sequence 34, Appl
9	38	50.7	79	9	US-09-796-692-2429	Sequence 2429, Ap
10	38	50.7	269	9	US-10-027-806-24	Sequence 24, Appl
11	38	50.7	269	9	US-10-034-623-24	Sequence 24, Appl
12	38	50.7	273	9	US-10-027-806-56	Sequence 56, Appl
13	38	50.7	273	9	US-10-034-623-56	Sequence 56, Appl
14	38	50.7	684	10	US-09-765-298A-18	Sequence 18, Appl
15	37	49.3	158	9	US-09-887-593-2	Sequence 2, Appli
16	37	49.3	194	10	US-09-764-898-285	Sequence 285, App
17	37	49.3	293	10	US-09-764-898-213	Sequence 213, App
18	37	49.3	405	12	US-10-060-333-2	Sequence 2, Appli
19	37	49.3	447	10	US-09-888-615-109	Sequence 109, App
20	37	49.3	533	9	US-10-041-406-2	Sequence 2, Appli
21	36	48.0	49	10	US-09-864-761-40695	Sequence 40695, A
22	36	48.0	452	10	US-09-925-297-683	Sequence 683, App
23	36	48.0	468	10	US-09-768-826-40	Sequence 40, Appl
24	36	48.0	494	10	US-09-833-790-234	Sequence 234, App
25	36	48.0	551	10	US-09-897-214-8	Sequence 8, Appli
26	36	48.0	565	10	US-09-768-826-58	Sequence 58, Appl
27	36	48.0	763	10	US-09-815-242-13643	Sequence 13643, A
28	35.5	47.3	221	10	US-09-764-864-1484	Sequence 1484, Ap
29	35.5	47.3	284	10	US-09-925-300-1322	Sequence 1322, Ap
30	35.5	47.3	513	10	US-09-764-864-1061	Sequence 1061, Ap
31	35.5	47.3	910	9	US-09-908-153B-40	Sequence 40, Appl
32	35.5	47.3	922	9	US-09-908-153B-42	Sequence 42, Appl
33	35	46.7	36	9	US-10-016-634A-161	Sequence 161, App
34	35	46.7	320	9	US-09-738-626-6378	Sequence 6378, Ap
35	35	46.7	523	10	US-09-815-242-11918	Sequence 11918, A
36	34.5	46.0	711	10	US-09-976-165-10	Sequence 10, Appl
37	34.5	46.0	711	10	US-09-828-648-2	Sequence 2, Appli
38	34	45.3	26	10	US-09-864-761-41575	Sequence 41575, A
39	34	45.3	36	10	US-09-864-761-34767	Sequence 34767, A
40	34	45.3	94	10	US-09-826-463-3	GENERAL INFORMA
41	34	45.3	100	10	US-09-893-238-126	Sequence 126, App
42	34	45.3	112	10	US-09-867-550-1070	Sequence 1070, Ap
43	34	45.3	322	10	US-09-741-669-404	Sequence 404, App
44	34	45.3	339	9	US-10-036-041-80	Sequence 80, Appl
45	34	45.3	339	10	US-09-729-674-138	Sequence 138, App

ALIGNMENTS

RESULT 1
US-10-007-968-15
; Sequence 15, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/10/007,968
; CURRENT FILING DATE: 2001-12-13
; PRIOR APPLICATION NUMBER: 09/740,211
; PRIOR FILING DATE: 2000-12-18
; PRIOR APPLICATION NUMBER: 60/125,974
; PRIOR FILING DATE: 1999-03-24
; PRIOR APPLICATION NUMBER: 60/104,994
; PRIOR FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 15
; LENGTH: 14
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-007-968-15

Query Match 100.0%; Score 75; DB 9; Length 14;
Best Local Similarity 100.0%; Pred. No. 8.8e-07;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db       1 SFSQNPPVLKRHQR 14
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RESULT 2
US-09-740-211-15
; Sequence 15, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:

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;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 15
;   LENGTH: 14
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-15

  Query Match             100.0%;  Score 75;  DB 10;  Length 14;
  Best Local Similarity   100.0%;  Pred. No. 8.8e-07;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 SFSQNPPVLKRHQR 14
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Db      1 SFSQNPPVLKRHQR 14


RESULT 3
US-10-006-091-1
; Sequence 1, Application US/10006091
; Patent No. US20020102730A1
; GENERAL INFORMATION:
;  APPLICANT: Cho, Myung-Sam
;  APPLICANT:  Chan, Sham-Yuen
;  APPLICANT:  Kelsey, William
;  APPLICANT:  Yee, Helena
;  TITLE OF INVENTION: Expression System for Factor VIII
;  FILE REFERENCE: MSB-7255.1
;  CURRENT APPLICATION NUMBER: US/10/006,091
;  CURRENT FILING DATE:  2001-12-06
;  NUMBER OF SEQ ID NOS: 2
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 1438
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Derived from
;   OTHER INFORMATION: human factor VIII sequence
US-10-006-091-1

  Query Match             100.0%;  Score 75;  DB 12;  Length 1438;
  Best Local Similarity   100.0%;  Pred. No. 0.00012;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 SFSQNPPVLKRHQR 14
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Db    741 SFSQNPPVLKRHQR 754


RESULT 4
US-10-047-257-1
; Sequence 1, Application US/10047257
; Patent No. US20020115152A1
; GENERAL INFORMATION:
;  APPLICANT: Cho, Myung-Sam
;  APPLICANT:  Chan, Sham-Yuen
;  APPLICANT:  Kelsey, William
;  APPLICANT:  Yee, Helena
;  TITLE OF INVENTION: Expression System for Factor VIII
;  FILE REFERENCE: MSB-7255.2
;  CURRENT APPLICATION NUMBER: US/10/047,257
;  CURRENT FILING DATE:  2002-01-15
;  NUMBER OF SEQ ID NOS: 2
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 1438
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Derived from
;   OTHER INFORMATION: human factor VIII sequence
US-10-047-257-1

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  Best Local Similarity   100.0%;  Pred. No. 0.00012;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 SFSQNPPVLKRHQR 14
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Db    741 SFSQNPPVLKRHQR 754


RESULT 5
US-09-957-641-2
; Sequence 2, Application US/09957641
; Publication No. US20020182670A1
; GENERAL INFORMATION:
;  APPLICANT: Emory University
;  TITLE OF INVENTION: MODIFIED FACTOR VIII
;  FILE REFERENCE: 75-00
;  CURRENT APPLICATION NUMBER: US/09/957,641
;  CURRENT FILING DATE:  2001-09-16
;  PRIOR APPLICATION NUMBER: US 60/234047
;  PRIOR FILING DATE: 2000-09-19
;  PRIOR APPLICATION NUMBER: US 60/236460
;  PRIOR FILING DATE: 2000-09-29
;  NUMBER OF SEQ ID NOS: 18
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
;   LENGTH: 2332
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-957-641-2

  Query Match             86.7%;  Score 65;  DB 9;  Length 2332;
  Best Local Similarity   100.0%;  Pred. No. 0.0091;
  Matches   12;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      3 SQNPPVLKRHQR 14
          ||||||||||||
Db   1637 SQNPPVLKRHQR 1648


RESULT 6
US-10-095-718-2
; Sequence 2, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
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;  PRIOR FILING DATE: 2001-08-22
;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2
;   LENGTH: 1471
;   TYPE: PRT
;   ORGANISM: Homo sapiens B-domain deleted factor VIII
;   FEATURE:
;   OTHER INFORMATION: Homo sapiens BDD FVIII
US-10-095-718-2

  Query Match             77.3%;  Score 58;  DB 12;  Length 1471;
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  Matches   14;  Conservative    0;  Mismatches    0;  Indels   14;  Gaps    1;

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Db    760 SFSQNSRHPSTRQKQFNATPPVLKRHQR 787


RESULT 7
US-09-748-062-30
; Sequence 30, Application US/09748062
; Patent No. US20010016340A1
;   GENERAL INFORMATION:
;        APPLICANT: CHAPMAN, BARBARA
;                   BURKE, RAE LYN
;                   RASMUSSEN, MIRELLA EZBAN
;                   MIKKELSON, JAN MOLLER
;        TITLE OF INVENTION: PROTEIN COMPLEXES HAVING FACTOR VIII:C
;                            ACTIVITY AND PRODUCTION THEREOF
;        NUMBER OF SEQUENCES: 35
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: REED & ROBINS
;             STREET: 285 HAMILTON AVENUE, SUITE 200
;             CITY: PALO ALTO
;             STATE: CALIFORNIA
;             COUNTRY: UNITED STATES OF AMERICA
;             ZIP: 94301
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.25
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/748,062
;             FILING DATE: 22-Dec-2000
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 08/441,943
;             FILING DATE: <Unknown>
;             APPLICATION NUMBER: US 822,989
;             FILING DATE: 27-JAN-1986
;        ATTORNEY/AGENT INFORMATION:
;             NAME: BAROVSKY, KENNETH
;             REGISTRATION NUMBER: 36,442
;             REFERENCE/DOCKET NUMBER: 2300-0048.10
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (415) 327-3400
;             TELEFAX: (415) 327-3231
;   INFORMATION FOR SEQ ID NO: 30:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 34 amino acids
;             TYPE: amino acid
;             TOPOLOGY: linear
;        MOLECULE TYPE: protein
;        SEQUENCE DESCRIPTION: SEQ ID NO: 30:
US-09-748-062-30

  Query Match             73.3%;  Score 55;  DB 10;  Length 34;
  Best Local Similarity   41.2%;  Pred. No. 0.0046;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels   20;  Gaps    1;

Qy      1 SFSQN--------------------PPVLKRHQR 14
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Db      1 SFSQNSRHPSTRQKQFNATPPTPPTPPVLKRHQR 34


RESULT 8
US-09-748-062-34
; Sequence 34, Application US/09748062
; Patent No. US20010016340A1
;   GENERAL INFORMATION:
;        APPLICANT: CHAPMAN, BARBARA
;                   BURKE, RAE LYN
;                   RASMUSSEN, MIRELLA EZBAN
;                   MIKKELSON, JAN MOLLER
;        TITLE OF INVENTION: PROTEIN COMPLEXES HAVING FACTOR VIII:C
;                            ACTIVITY AND PRODUCTION THEREOF
;        NUMBER OF SEQUENCES: 35
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: REED & ROBINS
;             STREET: 285 HAMILTON AVENUE, SUITE 200
;             CITY: PALO ALTO
;             STATE: CALIFORNIA
;             COUNTRY: UNITED STATES OF AMERICA
;             ZIP: 94301
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.25
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/748,062
;             FILING DATE: 22-Dec-2000
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 08/441,943
;             FILING DATE: <Unknown>
;             APPLICATION NUMBER: US 822,989
;             FILING DATE: 27-JAN-1986
;        ATTORNEY/AGENT INFORMATION:
;             NAME: BAROVSKY, KENNETH
;             REGISTRATION NUMBER: 36,442
;             REFERENCE/DOCKET NUMBER: 2300-0048.10
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (415) 327-3400
;             TELEFAX: (415) 327-3231
;   INFORMATION FOR SEQ ID NO: 34:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 9 amino acids
;             TYPE: amino acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        MOLECULE TYPE: peptide
;        SEQUENCE DESCRIPTION: SEQ ID NO: 34:
US-09-748-062-34

  Query Match             66.7%;  Score 50;  DB 10;  Length 9;
  Best Local Similarity   100.0%;  Pred. No. 9.8e+04;
  Matches    9;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      6 PPVLKRHQR 14
          |||||||||
Db      1 PPVLKRHQR 9


RESULT 9
US-09-796-692-2429
; Sequence 2429, Application US/09796692
; Publication No. US20020198362A1
; GENERAL INFORMATION:
;  APPLICANT: Gaiger, Alexander
;  APPLICANT:  Algate, Paul A.
```

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;  APPLICANT:  Mannion, Jane
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DETECTION, DIAGNOSIS AND THERAPY
;  TITLE OF INVENTION:  HEMATOLOGICAL MALIGNANCIES
;  FILE REFERENCE: 2077.001200
;  CURRENT APPLICATION NUMBER: US/09/796,692
;  CURRENT FILING DATE:  2001-03-01
;  PRIOR APPLICATION NUMBER: 60/186,126
;  PRIOR FILING DATE: 2000-03-01
;  PRIOR APPLICATION NUMBER: 60/190,479
;  PRIOR FILING DATE: 2000-03-17
;  PRIOR APPLICATION NUMBER: 60/200,545
;  PRIOR FILING DATE: 2000-04-27
;  PRIOR APPLICATION NUMBER: 60/200,303
;  PRIOR FILING DATE: 2000-04-28
;  PRIOR APPLICATION NUMBER: 60/200,779
;  PRIOR FILING DATE: 2000-04-28
;  PRIOR APPLICATION NUMBER: 60/200,999
;  PRIOR FILING DATE: 2000-05-01
;  PRIOR APPLICATION NUMBER: 60/202,084
;  PRIOR FILING DATE: 2000-05-04
;  PRIOR APPLICATION NUMBER: 60/206,201
;  PRIOR FILING DATE: 2000-05-22
;  PRIOR APPLICATION NUMBER: 60/218,950
;  PRIOR FILING DATE: 2000-07-14
;  PRIOR APPLICATION NUMBER: 60/222,903
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: 60/223,416
;  PRIOR FILING DATE: 2000-08-04
;  PRIOR APPLICATION NUMBER: 60/223,378
;  PRIOR FILING DATE: 2000-08-07
;  NUMBER OF SEQ ID NOS: 9597
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2429
;   LENGTH: 79
;   TYPE: PRT
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: variant
;   LOCATION: (1)...(79)
;   OTHER INFORMATION: Xaa = Any amino acid
US-09-796-692-2429

  Query Match             50.7%;  Score 38;  DB 9;  Length 79;
  Best Local Similarity   77.8%;  Pred. No. 7.4;
  Matches    7;  Conservative    1;  Mismatches    1;  Indels    0;  Gaps    0;

Qy       1 SFSQNPPVL 9
           ||| |||:|
Db      43 SFSPNPPLL 51


RESULT 10
US-10-027-806-24
; Sequence 24, Application US/10027806
; Patent No. US20020160476A1
; GENERAL INFORMATION:
;  APPLICANT: Swanson, Ronald V.
;  APPLICANT:  Feldman, Robert A.
;  APPLICANT:  Schleper, Christa
;  TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
;  FILE REFERENCE: DCORP.002A
;  CURRENT APPLICATION NUMBER: US/10/027,806
;  CURRENT FILING DATE:  2001-12-21
;  PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/408,020
;  PRIOR FILING DATE: EARLIER FILING DATE: 1999-09-29
;  NUMBER OF SEQ ID NOS: 123
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 24
;   LENGTH: 269
;   TYPE: PRT
;   ORGANISM: Cenarchaeum symbiosum
US-10-027-806-24

  Query Match             50.7%;  Score 38;  DB 9;  Length 269;
  Best Local Similarity   63.6%;  Pred. No. 27;
  Matches    7;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy       4 QNPPVLKRHQR 14
           |||||::|  |
Db      50 QNPPVIQRGGR 60


RESULT 11
US-10-034-623-24
; Sequence 24, Application US/10034623
; Publication No. US20020198365A1
; GENERAL INFORMATION:
;  APPLICANT: Swanson, Ronald V.
;  APPLICANT:  Feldman, Robert A.
;  APPLICANT:  Schleper, Christa
;  TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
;  FILE REFERENCE: DCORP.002A
;  CURRENT APPLICATION NUMBER: US/10/034,623
;  CURRENT FILING DATE:  2001-12-21
;  PRIOR APPLICATION NUMBER: 09/408,020
;  PRIOR FILING DATE: 1999-09-29
;  PRIOR APPLICATION NUMBER: 60/102,294
;  PRIOR FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 123
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 24
;   LENGTH: 269
;   TYPE: PRT
;   ORGANISM: Cenarchaeum symbiosum
US-10-034-623-24

  Query Match             50.7%;  Score 38;  DB 9;  Length 269;
  Best Local Similarity   63.6%;  Pred. No. 27;
  Matches    7;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy       4 QNPPVLKRHQR 14
           |||||::|  |
Db      50 QNPPVIQRGGR 60


RESULT 12
US-10-027-806-56
; Sequence 56, Application US/10027806
; Patent No. US20020160476A1
; GENERAL INFORMATION:
;  APPLICANT: Swanson, Ronald V.
;  APPLICANT:  Feldman, Robert A.
;  APPLICANT:  Schleper, Christa
;  TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
;  FILE REFERENCE: DCORP.002A
;  CURRENT APPLICATION NUMBER: US/10/027,806
;  CURRENT FILING DATE:  2001-12-21
;  PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/408,020
;  PRIOR FILING DATE: EARLIER FILING DATE: 1999-09-29
;  NUMBER OF SEQ ID NOS: 123
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 56
;   LENGTH: 273
;   TYPE: PRT
;   ORGANISM: Cenarchaeum symbiosum
US-10-027-806-56

  Query Match             50.7%;  Score 38;  DB 9;  Length 273;
  Best Local Similarity   63.6%;  Pred. No. 28;
  Matches    7;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy       4 QNPPVLKRHQR 14
           |||||::|  |
Db      54 QNPPVIQRGGR 64
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RESULT 13
US-10-034-623-56
; Sequence 56, Application US/10034623
; Publication No. US20020198365A1
; GENERAL INFORMATION:
;  APPLICANT: Swanson, Ronald V.
;  APPLICANT:  Feldman, Robert A.
;  APPLICANT:  Schleper, Christa
;  TITLE OF INVENTION: NUCLEIC ACIDS AND PROTEINS FROM CENARCHAEUM SYMBIOSUM
;  FILE REFERENCE: DCORP.002A
;  CURRENT APPLICATION NUMBER: US/10/034,623
;  CURRENT FILING DATE:  2001-12-21
;  PRIOR APPLICATION NUMBER: 09/408,020
;  PRIOR FILING DATE: 1999-09-29
;  PRIOR APPLICATION NUMBER: 60/102,294
;  PRIOR FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 123
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 56
;   LENGTH: 273
;   TYPE: PRT
;   ORGANISM: Cenarchaeum symbiosum
US-10-034-623-56

  Query Match             50.7%;  Score 38;  DB 9;  Length 273;
  Best Local Similarity   63.6%;  Pred. No. 28;
  Matches    7;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy       4 QNPPVLKRHQR 14
           |||||::|  |
Db      54 QNPPVIQRGGR 64


RESULT 14
US-09-765-298A-18
; Sequence 18, Application US/09765298A
; Patent No. US20020137017A1
; GENERAL INFORMATION:
;  APPLICANT: ARONHEIM, AMI
;  TITLE OF INVENTION: METHOD FOR DETECTION PROTEIN-PROTEIN INTERACTIONS AND A KIT THERE
;  FILE REFERENCE: 108387.01
;  CURRENT APPLICATION NUMBER: US/09/765,298A
;  CURRENT FILING DATE:  2001-01-22
;  PRIOR APPLICATION NUMBER: IL 125456
;  PRIOR FILING DATE: 1998-07-22
;  PRIOR APPLICATION NUMBER: IL 128017
;  PRIOR FILING DATE: 1999-01-12
;  NUMBER OF SEQ ID NOS: 31
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
;   LENGTH: 684
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-765-298A-18

  Query Match             50.7%;  Score 38;  DB 10;  Length 684;
  Best Local Similarity   87.5%;  Pred. No. 73;
  Matches    7;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy       5 NPPVLKRH 12
           || |||||
Db     295 NPEVLKRH 302


RESULT 15
US-09-887-593-2
; Sequence 2, Application US/09887593
; Patent No. US20020161212A1
; GENERAL INFORMATION:
;  APPLICANT: Afar, Daniel E.
;  APPLICANT:  Hubert, Rene S.
;  APPLICANT:  Leong, Kahan
;  APPLICANT:  Raitano, Arthur B.
;  APPLICANT:  Saffran, Douglas C.
;  APPLICANT:  Jakobovits, Aya
;  TITLE OF INVENTION: BPC-1: A SECRETED BRAIN-SPECIFIC PROTEIN EXPRESSED AND
;  TITLE OF INVENTION:  SECRETED BY PROSTATE AND BLADDER CANCER CELLS
;  FILE REFERENCE: 1703-017.US1
;  CURRENT APPLICATION NUMBER: US/09/887,593
;  CURRENT FILING DATE:  2001-06-21
;  PRIOR APPLICATION NUMBER: 09/374,135
;  PRIOR FILING DATE: 1999-08-10
;  PRIOR APPLICATION NUMBER: 60/095,982
;  PRIOR FILING DATE: 1998-08-10
;  NUMBER OF SEQ ID NOS: 20
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
;   LENGTH: 158
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-887-593-2

  Query Match             49.3%;  Score 37;  DB 9;  Length 158;
  Best Local Similarity   63.6%;  Pred. No. 23;
  Matches    7;  Conservative    1;  Mismatches    3;  Indels    0;  Gaps    0;

Qy       4 QNPPVLKRHQR 14
           |||||:|   |
Db     121 QNPPVIKSSGR 131


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Job time : 19 secs
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GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: January 4, 2003, 04:58:21 ; Search time 47.5 Seconds
(without alignments)
6462.809 Million cell updates/sec

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Perfect score: 1001
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

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Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1001	100.0	4629	2	US-08-484-891-7	Sequence 7, Appli
2	1001	100.0	4670	3	US-08-717-294-41	Sequence 41, Appl
3	1001	100.0	4999	4	US-09-470-618-14	Sequence 14, Appl
4	1001	100.0	4999	4	US-09-364-862-14	Sequence 14, Appl
5	1001	100.0	5035	2	US-08-882-083-1	Sequence 1, Appli
6	1001	100.0	5035	2	US-08-558-107-1	Sequence 1, Appli
7	1001	100.0	5035	3	US-09-243-539-1	Sequence 1, Appli
8	1001	100.0	7056	1	US-08-121-202-1	Sequence 1, Appli
9	1001	100.0	8241	6	5171844-1	Patent No. 5171844
10	1001	100.0	8967	1	US-08-366-851A-1	Sequence 1, Appli
11	1001	100.0	9009	1	US-07-864-004B-3	Sequence 3, Appli
12	1001	100.0	9009	1	US-08-251-937A-3	Sequence 3, Appli
13	1001	100.0	9009	1	US-08-212-133A-1	Sequence 1, Appli
14	1001	100.0	9009	1	US-08-474-503-1	Sequence 1, Appli
15	1001	100.0	9009	2	US-08-670-707A-1	Sequence 1, Appli
16	1001	100.0	9009	4	US-09-037-601-1	Sequence 1, Appli
17	1001	100.0	9009	4	US-09-315-179-1	Sequence 1, Appli
18	1001	100.0	9009	4	US-09-523-656-1	Sequence 1, Appli
19	1001	100.0	9009	5	PCT-US93-03275-3	Sequence 3, Appli
20	1001	100.0	9009	5	PCT-US94-13200-1	Sequence 1, Appli
21	1001	100.0	9354	1	US-08-683-839B-2	Sequence 2, Appli
22	1001	100.0	11933	4	US-09-470-618-13	Sequence 13, Appl
23	1001	100.0	11933	4	US-09-364-862-13	Sequence 13, Appl
24	997.8	99.7	6999	1	US-08-276-594A-1	Sequence 1, Appli
25	858.6	85.8	7032	4	US-09-324-867-1	Sequence 1, Appli
26	831.4	83.1	7493	1	US-08-212-133A-7	Sequence 7, Appli
27	831.4	83.1	7493	1	US-08-474-503-5	Sequence 5, Appli
28	831.4	83.1	7493	2	US-08-670-707A-5	Sequence 5, Appli
29	831.4	83.1	7493	4	US-09-037-601-5	Sequence 5, Appli
30	831.4	83.1	7493	4	US-09-315-179-5	Sequence 5, Appli
31	831.4	83.1	7493	5	PCT-US94-13200-5	Sequence 5, Appli
32	810.6	81.0	4404	4	US-09-523-656-37	Sequence 37, Appl
33	809	80.8	4334	2	US-08-670-707A-38	Sequence 38, Appl
34	809	80.8	4334	4	US-09-037-601-38	Sequence 38, Appl
35	809	80.8	4334	4	US-09-315-179-38	Sequence 38, Appl
36	809	80.8	6402	2	US-08-670-707A-36	Sequence 36, Appl
37	809	80.8	6402	4	US-09-037-601-36	Sequence 36, Appl
38	809	80.8	6402	4	US-09-315-179-36	Sequence 36, Appl
39	809	80.8	6402	4	US-09-523-656-29	Sequence 29, Appl
40	612.2	61.2	4451	3	US-08-717-294-42	Sequence 42, Appl
41	147.6	14.7	6909	2	US-08-804-196-1	Sequence 1, Appli
42	147.6	14.7	6909	2	US-08-658-340-1	Sequence 1, Appli
43	147.6	14.7	6909	3	US-08-746-111-26	Sequence 26, Appl
44	140	14.0	6585	3	US-08-746-111-4	Sequence 4, Appli
45	137.4	13.7	2012	4	US-09-149-476-132	Sequence 132, App

ALIGNMENTS

RESULT 1
US-08-484-891-7
; Sequence 7, Application US/08484891
; Patent No. 5935935
; GENERAL INFORMATION:
; APPLICANT: Connelly, Sheila
; APPLICANT: Kaleko, Michael
; APPLICANT: Smith, Theodore
; TITLE OF INVENTION: Adenoviral Vectors for
; TITLE OF INVENTION: Treatment of Hemophilia
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carella, Byrne, Bain, Gilfillan,
; ADDRESSEE: Cecchi, Stewart & Olstein
; STREET: 6 Becker Farm Road
; CITY: Roseland
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 inch diskette
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WordPerfect 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/484,891
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/218,335
; FILING DATE: 25-MAR-1994
; APPLICATION NUMBER: 08/074,920
; FILING DATE: 10-JUN-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Olstein, Elliot M.
; REGISTRATION NUMBER: 24,025
; REFERENCE/DOCKET NUMBER: 271010-273
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 201-994-1700
; TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4629 bases
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA primer
; FEATURE:
; NAME/KEY: Factor VIII cDNA with
; NAME/KEY: B domain deleted

US-08-484-891-7

Query Match 100.0%; Score 1001; DB 2; Length 4629;
Best Local Similarity 100.0%; Pred. No. 5.6e-310;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    2652 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 2711

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2712 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 2771

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Db    2772 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 2831

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2832 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 2891

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2892 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 2951

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2952 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3011

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3012 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3071

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3072 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3131

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3132 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3191

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3192 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3251

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3252 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3311

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3312 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3371

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3372 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3431

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3432 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3491

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3492 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3551

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3552 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3611

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    3612 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 3652
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RESULT 2
US-08-717-294-41
; Sequence 41, Application US/08717294
; Patent No. 6114148
; GENERAL INFORMATION:
; APPLICANT: SEED, BRIAN
; APPLICANT: HAAS, JURGEN
; TITLE OF INVENTION: HIGH LEVEL EXPRESSION OF
; TITLE OF INVENTION: PROTEINS
; NUMBER OF SEQUENCES: 110
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Clark & Elbing LLP
; STREET: 176 Federal Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/717,294
; FILING DATE: 20-SEP-1996
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Elbing, Karen L
; REGISTRATION NUMBER: 35,238
; REFERENCE/DOCKET NUMBER: 00786/345001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-428-0200
; TELEFAX: 617-428-7045
; TELEX:
; INFORMATION FOR SEQ ID NO: 41:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4670 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-717-294-41

Query Match 100.0%; Score 1001; DB 3; Length 4670;
Best Local Similarity 100.0%; Pred. No. 5.6e-310;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2678 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 2737

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2738 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 2797

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2798 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 2857

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2858 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 2917

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2918 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 2977

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2978 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3037

Qy    361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3038 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3097

Qy    421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3098 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3157

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3158 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3217

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3218 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3277

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3278 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3337

Qy    661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3338 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3397

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3398 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3457

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3458 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3517

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3518 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3577

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3578 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3637

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
        |||||||||||||||||||||||||||||||||||||||||
Db   3638 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 3678


RESULT 3
US-09-470-618-14
; Sequence 14, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION: by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/470,618
;  CURRENT FILING DATE: 1999-12-22
;  EARLIER APPLICATION NUMBER: 09/364,862
;  EARLIER FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-470-618-14

  Query Match           100.0%;  Score 1001;  DB 4;  Length 4999;
  Best Local Similarity  100.0%;  Pred. No. 5.8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3025 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3084

Qy     61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3085 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3144

Qy    121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3145 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3204

Qy    181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3205 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3264

Qy    241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3265 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3324

Qy    301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3325 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3384

Qy    361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3385 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3444

Qy    421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3445 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3504

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3505 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3564

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3565 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3624

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3625 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3684

Qy    661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3685 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3744

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3745 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3804

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3805 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3864

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3865 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3924

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3925 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3984

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
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Db   3985 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4025


RESULT 4
US-09-364-862-14
; Sequence 14, Application US/09364862
; Patent No. 6221349
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: ADENO-ASSOCIATED VECTORS FOR EXPRESSION OF FACTOR VIII
;  TITLE OF INVENTION: BY TARGET
;  TITLE OF INVENTION: CELLS
;  FILE REFERENCE: AVIGEN-03743
;  CURRENT APPLICATION NUMBER: US/09/364,862
;  CURRENT FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 14
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-364-862-14

  Query Match            100.0%;  Score 1001;  DB 4;  Length 4999;
  Best Local Similarity  100.0%;  Pred. No. 5.8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    3025 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3084

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3085 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3144

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3145 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3204

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3205 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3264

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3265 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3324

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3325 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3384

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3385 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3444

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3445 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3504

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3505 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3564

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3565 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3624

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3625 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3684

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3685 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3744

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3745 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3804

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3805 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3864

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3865 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3924

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3925 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3984

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          ||||||||||||||||||||||||||||||||||||||||||
Db    3985 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4025


RESULT 5
US-08-882-083-1
; Sequence 1, Application US/08882083
; Patent No. 5869292
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/882,083
;      FILING DATE:
;      CLASSIFICATION:  514
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
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;      LOCATION:  35..5017
US-08-882-083-1

  Query Match            100.0%;  Score 1001;  DB 2;  Length 5035;
  Best Local Similarity   100.0%;  Pred. No. 5.8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3256 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3315

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3316 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3375

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3376 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3435

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3436 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3495

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
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Db    3496 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3555

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3556 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3615

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3616 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3675

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3676 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3735

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3736 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3795

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3796 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3855

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3856 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3915

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3916 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3975

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3976 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 4035

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4036 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 4095

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4096 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 4155

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4156 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 4215

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db    4216 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4256


RESULT 6
US-08-558-107-1
; Sequence 1, Application US/08558107
; Patent No. 5910481
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/558,107
;      FILING DATE:  13-NOV-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  35..5017
US-08-558-107-1

  Query Match            100.0%;  Score 1001;  DB 2;  Length 5035;
  Best Local Similarity   100.0%;  Pred. No. 5.8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3256 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3315

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3316 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3375

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3376 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3435

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3436 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3495

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3496 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3555

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3556 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3615
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Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
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Db    3616 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3675

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3676 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3735

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3736 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3795

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3796 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3855

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3856 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3915

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3916 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3975

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3976 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 4035

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
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Db    4036 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 4095

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
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Db    4096 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 4155

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
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Db    4156 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 4215

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
           ||||||||||||||||||||||||||||||||||||||||
Db    4216 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4256
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RESULT 7
US-09-243-539-1
; Sequence 1, Application US/09243539
; Patent No. 6130203
;  GENERAL INFORMATION:
;    APPLICANT:  VOORBERG, Johannes J.
;    TITLE OF INVENTION:  HYBRID PROTEINS WITH MODIFIED ACTIVITY
;    NUMBER OF SEQUENCES:  17
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Foley & Lardner
;      STREET:  3000 K Street, N.W., Suite 500
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20007-5109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/243,539
;      FILING DATE:
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/558,107
;      FILING DATE:  13-NOV-1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ISACSON, John P.
;      REGISTRATION NUMBER:  33,715
;      REFERENCE/DOCKET NUMBER:  30472/212
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)672-5300
;      TELEFAX:  (202)672-5399
;      TELEX:  904136
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5035 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  35..5017
US-09-243-539-1

  Query Match           100.0%;  Score 1001;  DB 3;  Length 5035;
  Best Local Similarity  100.0%;  Pred. No. 5.8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    3256 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3315

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
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Db    3316 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3375

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Db    3376 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3435

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3436 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3495

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
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Db    3496 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3555

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
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Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3676 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3735

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
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Db    3736 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3795

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
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Db    3796 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3855

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
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Db    3856 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3915

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
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Db    3916 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3975

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Db   3976 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 4035

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4036 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 4095

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4096 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 4155

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
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Db   4156 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 4215

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   4216 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4256


RESULT 8
US-08-121-202-1
; Sequence 1, Application US/08121202
; Patent No. 5563045
;  GENERAL INFORMATION:
;    APPLICANT:  Pittman, Debra
;    APPLICANT:  Rehemtulla, Alnawaz
;    APPLICANT:  Wozney, John M.
;    APPLICANT:  Kaufman, Randal J.
;    TITLE OF INVENTION:  CHIMERIC PROCOAGULANT PROTEINS
;    NUMBER OF SEQUENCES:  27
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Legal Affairs, Genetics Institute, Inc.
;      STREET:  87 CambridgePark Drive
;      CITY:  Cambridge
;      STATE:  MA
;      COUNTRY:  USA
;      ZIP:  02140
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/121,202
;      FILING DATE:  14-SEP-1993
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Meinert, M. C.
;      REGISTRATION NUMBER:  31,544
;      REFERENCE/DOCKET NUMBER:  GI 5195A
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)876-1210 X8574
;      TELEFAX:  (617) 876-5851
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  7056 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..7053
US-08-121-202-1

  Query Match            100.0%;  Score 1001;  DB 1;  Length 7056;
  Best Local Similarity  100.0%;  Pred. No. 7e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db   5292 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5351

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Db   5352 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5411

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Db   5412 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5471

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Db   5472 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5531

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5532 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5591

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5592 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5651

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5652 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5711

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
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Db   5712 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5771

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5772 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5831

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5832 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 5891

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5892 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 5951

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5952 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6011

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6012 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6071

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6072 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6131

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6132 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6191

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6192 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6251

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
           |||||||||||||||||||||||||||||||||||||||||
Db   6252 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6292


RESULT 9
5171844-1
;Patent No. 5171844
;    APPLICANT: VAN OOYEN, ALBERT J.J.;PANNEKOEK, HANS;VERBEET,
;MARTINUS P.;VAN LEEN, ROBERT W.
;    TITLE OF INVENTION: PROTEINS WITH FACTOR VIII ACTIVITY
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;PROCESS FOR THEIR PREPARATION USING GENETICALLY-ENGINEERED CELLS
;AND PHARMACEUTICAL COMPOSITIONS CONTAINING THEM
;     NUMBER OF SEQUENCES: 12
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/07/205,226
;       FILING DATE: 10-JUN-1988
;SEQ ID NO:1:
;       LENGTH: 8241
5171844-1

  Query Match             100.0%;  Score 1001;  DB 6;  Length 8241;
  Best Local Similarity   100.0%;  Pred. No. 7.6e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5298 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5357

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5358 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5417

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5418 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5477

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5478 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5537

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5538 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5597

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5598 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5657

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5658 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5717

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5718 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5777

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5778 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5837

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5838 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 5897

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5898 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 5957

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5958 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6017

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6018 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6077

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6078 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6137

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6138 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6197

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6198 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6257

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          ||||||||||||||||||||||||||||||||||||||||||
Db    6258 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6298


RESULT 10
US-08-366-851A-1
; Sequence 1, Application US/08366851A
; Patent No. 5681746
;  GENERAL INFORMATION:
;    APPLICANT:  Bodner, Mordechai
;    APPLICANT:  De Polo, Nicolas J.
;    APPLICANT:  Hsu, David Chi-Tang
;    APPLICANT:  Chang, Steven
;    TITLE OF INVENTION:  Retroviral Delivery of Full Length Factor VIII
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Viagene, Inc.
;      STREET:  11055 Roselle Street
;      CITY:  San Diego
;      STATE:  California
;      COUNTRY:  U.S.A.
;      ZIP:  92121
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/366,851A
;      FILING DATE:
;      CLASSIFICATION:  514
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Chambers, Daniel M.
;      REGISTRATION NUMBER:  34,561
;      REFERENCE/DOCKET NUMBER:  930049.438
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (619) 452-1288
;      TELEFAX:  (619) 452-2616
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  8967 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  110..7165
US-08-366-851A-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 8967;
  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5401 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5460

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5461 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5520

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5521 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5580

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Db   5581 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5640

Qy    241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5641 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5700

Qy    301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5701 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5760

Qy    361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5761 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5820

Qy    421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5821 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5880

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5881 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5940

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5941 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6000

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6001 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6060

Qy    661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6061 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6120

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6121 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6180

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6181 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6240

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6241 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6300

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6301 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6360

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   6361 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6401


RESULT 11
US-07-864-004B-3
; Sequence 3, Application US/07864004B
; Patent No. 5364771
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  6
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/864,004B
;      FILING DATE:  07 APRIL 1992
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU106
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6508
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5001 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2
;      OTHER INFORMATION:  domain"
US-07-864-004B-3

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5442 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5501

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5502 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5561

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5562 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5621

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5622 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5681

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5682 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5741

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5742 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5801

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5802 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5861

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
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Db    5862 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5921

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
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Db    5922 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5981

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5982 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6041

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6042 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6101

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6102 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6161

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6162 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6221

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6222 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6281

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6282 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6341

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6342 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6401

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
                ||||||||||||||||||||||||||||||||||||||||
Db    6402 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6442


RESULT 12
US-08-251-937A-3
; Sequence 3, Application US/08251937A
; Patent No. 5583209
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Porcine Factor VIII
;    NUMBER OF SEQUENCES:  10
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/251,937A
;      FILING DATE:  31-MAY-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106DIV
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6367
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FRAGMENT TYPE:  N-terminal
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5001 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2
;      OTHER INFORMATION:  domain"
US-08-251-937A-3

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    5442 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5501

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
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Db    5502 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5561

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Db    5562 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5621

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Db    5622 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5681

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
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Db    5682 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5741

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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Db    5742 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5801

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
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Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
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Db    5922 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5981

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Db    5982 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6041

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db   6042 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6101

Qy    661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
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Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6162 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6221

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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   6222 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6281

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
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Db   6282 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6341

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
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Db   6342 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6401

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RESULT 13
US-08-212-133A-1
; Sequence 1, Application US/08212133A
; Patent No. 5663060
;  GENERAL INFORMATION:
;    APPLICANT:  Lollar, John S.
;    APPLICANT:  Runge, Marschall S.
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  100 Peachtree Street
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30303
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/212,133A
;      FILING DATE:  March 11, 1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/864,004
;      FILING DATE:  07-APR-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pabst, Patrea L.
;      REGISTRATION NUMBER:  31,284
;      REFERENCE/DOCKET NUMBER:  EMU/76677
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-572-6508
;      TELEFAX:  404-572-6555
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5125 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2 domain."
;    FEATURE:
;      NAME/KEY:  Domain
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /note= "cDNA encoding human factor
;      OTHER INFORMATION:  VIII."
US-08-212-133A-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    5442 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5501

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
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Db    5502 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5561

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
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Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
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Db    5862 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5921

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
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Db    5982 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6041

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6042 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6101

Qy     661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
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Db    6102 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6161

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
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Db    6162 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6221
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Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
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Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
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RESULT 14
US-08-474-503-1
; Sequence 1, Application US/08474503
; Patent No. 5744446
;  GENERAL INFORMATION:
;    APPLICANT:  Emory University
;    TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kilpatrick & Cody
;      STREET:  1100 Peachtree Street, Suite 2800
;      CITY:  Atlanta
;      STATE:  Georgia
;      COUNTRY:  US
;      ZIP:  30309
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/474,503
;      FILING DATE:  07-JUN-1995
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Pratt, John S.
;      REGISTRATION NUMBER:  29,476
;      REFERENCE/DOCKET NUMBER:  EMU106CIP(3)
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  404-815-6500
;      TELEFAX:  404-815-6555
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  9009 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapien
;      TISSUE TYPE:  Liver
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  5125 . . . 7053
;      OTHER INFORMATION:  /note= "Equivalent to the A3-C1-C2
;      OTHER INFORMATION:  domain"
;    FEATURE:
;      NAME/KEY:  misc_feature (Domain Structure)
;      LOCATION:  1 . . . 2277
;      OTHER INFORMATION:  /note= "Equivalent to the A1-A2 domain."
;    FEATURE:
;      NAME/KEY:  Domain
;      LOCATION:  1..2277
;      OTHER INFORMATION:  /note= "cDNA encoding human factor
;      OTHER INFORMATION:  VIII."
US-08-474-503-1

  Query Match             100.0%;  Score 1001;  DB 1;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
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Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
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Db    5802 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5861

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
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Db    5862 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5921

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
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Db    5922 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5981

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
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Db    5982 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6041

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
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Db    6042 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6101

Qy     661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6102 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6161

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6162 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6221

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    6222 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6281

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
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Db    6282 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6341

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
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Db   6402 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6442


RESULT 15
US-08-670-707A-1
; Sequence 1, Application US/08670707A
; Patent No. 5859204
; GENERAL INFORMATION:
;   APPLICANT:  Lollar, John S.
;   TITLE OF INVENTION:  Hybrid Human/Animal Factor VIII
;   NUMBER OF SEQUENCES:  40
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE:  Greenlee, Winner and Sullivan, P.C.
;     STREET:  5370 Manhattan Circle Suite 201
;     CITY:  Boulder
;     STATE:  Colorado
;     COUNTRY:  USA
;     ZIP:  80303
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Floppy disk
;     COMPUTER:  IBM PC compatible
;     OPERATING SYSTEM:  PC-DOS/MS-DOS
;     SOFTWARE:  PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER:  US/08/670,707A
;     FILING DATE:  26-JUN-1996
;     CLASSIFICATION:  435
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  WO PCT/US94/13200
;     FILING DATE:  15-NOV-1994
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  US 08/212,133
;     FILING DATE:  11-MAR-1994
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  US 07/864,004
;     FILING DATE:  07-APR-1992
;   ATTORNEY/AGENT INFORMATION:
;     NAME:  Greenlee, Lorance L.
;     REGISTRATION NUMBER:  27,894
;     REFERENCE/DOCKET NUMBER:  75-95F
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  303/499-8080
;     TELEFAX:  303/499-8089
; INFORMATION FOR SEQ ID NO:  1:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH:  9009 base pairs
;     TYPE:  nucleic acid
;     STRANDEDNESS:  double
;     TOPOLOGY:  not relevant
;   MOLECULE TYPE:  cDNA to mRNA
;   HYPOTHETICAL:  NO
;   ANTI-SENSE:  NO
;   ORIGINAL SOURCE:
;     ORGANISM:  Homo sapiens
;     TISSUE TYPE:  Liver
;   FEATURE:
;     NAME/KEY:  misc_feature
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;     OTHER INFORMATION:  /product= "Domain Structure"
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US-08-670-707A-1

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  Best Local Similarity   100.0%;  Pred. No. 8e-310;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
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Db    5442 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5501

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
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Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
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Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
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Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
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Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
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Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
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Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
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Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
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Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
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Search completed: January  4, 2003, 05:20:02
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Job time : 209.5 secs

GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: January 4, 2003, 04:59:31 ; Search time 53.5 Seconds
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Perfect score: 1001
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 381593 seqs, 216252194 residues

Total number of hits satisfying chosen parameters: 763186

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2		1001	100.0	4999	9	US-10-007-968-14	Sequence 14, Appl
3		1001	100.0	4999	10	US-09-740-211-14	Sequence 14, Appl
4		1001	100.0	7944	12	US-10-095-718-1	Sequence 1, Appli
5		1001	100.0	9009	9	US-09-957-641-1	Sequence 1, Appli
6		1001	100.0	11933	9	US-10-007-968-13	Sequence 13, Appl
7		1001	100.0	11933	10	US-09-740-211-13	Sequence 13, Appl
8		857	85.6	7914	12	US-10-095-718-3	Sequence 3, Appli
9		147.6	14.7	6909	10	US-09-880-107-2275	Sequence 2275, Ap
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c 21		44.6	4.5	404	10	US-09-778-320-132	Sequence 132, App
c 22		44.6	4.5	404	10	US-09-910-689-132	Sequence 132, App
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39		34	3.4	197997	10	US-09-822-246-3	Sequence 3, Appli
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c 42		33.6	3.4	1503841	9	US-09-946-807-1	Sequence 1, Appli
c 43		33.6	3.4	1503841	10	US-09-795-668-1	Sequence 1, Appli
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c 45		33	3.3	259	10	US-09-777-564-1384	Sequence 1384, Ap

ALIGNMENTS

RESULT 1
US-09-150-811-7
; GENERAL INFORMATION:
; APPLICANT: Connelly, Sheila
; Kaleko, Michael
; Smith, Theodore
; TITLE OF INVENTION: Adenoviral Vectors for Treatment of Hemophilia
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carella, Byrne, Bain, Gilfillan, Cecchi, Stewart & Olste
; STREET: 6 Becker Farm Road
; CITY: Roseland
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 inch diskette
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WordPerfect 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/150,811
; FILING DATE: 10-Sep-1998
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/484,891
; FILING DATE: 07-JUN-1995
; APPLICATION NUMBER: 08/218,335
; FILING DATE: 25-MAR-1994
; APPLICATION NUMBER: 08/074,920
; FILING DATE: 10-JUN-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Olstein, Elliot M.
; REGISTRATION NUMBER: 24,025
; REFERENCE/DOCKET NUMBER: 271010-440
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 973-994-1700
; TELEFAX: 973-994-1744
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-150-811-7

Query Match 100.0%; Score 1001; DB 10; Length 4629;

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RESULT 2
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; Sequence 14, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
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;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-14

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Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3565 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3624

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db   3625 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3684

Qy    661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3685 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3744

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3745 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3804

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3805 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3864

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3865 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3924

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3925 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3984

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   3985 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4025


RESULT 3
US-09-740-211-14
; Sequence 14, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
;   LENGTH: 4999
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-14

  Query Match             100.0%;  Score 1001;  DB 10;  Length 4999;
  Best Local Similarity   100.0%;  Pred. No. 2.6e-299;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3025 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3084

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3085 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3144

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3145 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3204

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3205 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3264

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3265 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3324

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3325 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3384

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3385 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3444

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3445 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3504

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3505 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3564

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3565 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3624

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3625 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3684

Qy     661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3685 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3744

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3745 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3804

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3805 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3864

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3865 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3924

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3925 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3984

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||
Db   3985 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4025


RESULT 4
US-10-095-718-1
; Sequence 1, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
;  PRIOR FILING DATE: 2001-08-22
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;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
;   LENGTH: 7944
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Plasmid pDLZ6 encoding Homo sapiens BDD FVIII
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (420)...(4835)
US-10-095-718-1

  Query Match             100.0%;  Score 1001;  DB 12;  Length 7944;
  Best Local Similarity   100.0%;  Pred. No. 3.4e-299;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3071 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3130

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3131 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3190

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3191 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3250

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3251 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3310

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3311 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3370

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3371 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3430

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3431 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3490

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3491 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3550

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3551 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3610

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3611 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3670

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3671 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3730

Qy     661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3731 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3790

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3791 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3850

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db    3851 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3910

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3911 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3970

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3971 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 4030

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          ||||||||||||||||||||||||||||||||||||||||||
Db    4031 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4071


RESULT 5
US-09-957-641-1
; Sequence 1, Application US/09957641
; Publication No. US20020182670A1
; GENERAL INFORMATION:
;  APPLICANT: Emory University
;  TITLE OF INVENTION: MODIFIED FACTOR VIII
;  FILE REFERENCE: 75-00
;  CURRENT APPLICATION NUMBER: US/09/957,641
;  CURRENT FILING DATE:  2001-09-16
;  PRIOR APPLICATION NUMBER: US 60/234047
;  PRIOR FILING DATE: 2000-09-19
;  PRIOR APPLICATION NUMBER: US 60/236460
;  PRIOR FILING DATE: 2000-09-29
;  NUMBER OF SEQ ID NOS: 18
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 9009
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (208)..(7203)
US-09-957-641-1

  Query Match             100.0%;  Score 1001;  DB 9;  Length 9009;
  Best Local Similarity   100.0%;  Pred. No. 3.7e-299;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5442 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 5501

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5502 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 5561

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5562 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 5621

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5622 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 5681

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5682 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 5741

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5742 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 5801

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    5802 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 5861

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
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Db   5862 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 5921

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
Db   5922 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 5981

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
Db   5982 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 6041

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
Db   6042 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 6101

Qy    661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
Db   6102 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 6161

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
Db   6162 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 6221

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
Db   6222 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 6281

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
Db   6282 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 6341

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
Db   6342 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 6401

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
Db   6402 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 6442


RESULT 6
US-10-007-968-13
; Sequence 13, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/10/007,968
;  CURRENT FILING DATE:  2001-12-13
;  PRIOR APPLICATION NUMBER: 09/740,211
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-10-007-968-13

  Query Match             100.0%;  Score 1001;  DB 9;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 4.3e-299;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy      1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
Db   3000 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3059

Qy     61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
Db   3060 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3119

Qy    121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
Db   3120 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3179

Qy    181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
Db   3180 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3239

Qy    241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
Db   3240 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3299

Qy    301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
Db   3300 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3359

Qy    361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
Db   3360 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3419

Qy    421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
Db   3420 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3479

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
Db   3480 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3539

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
Db   3540 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3599

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
Db   3600 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3659

Qy    661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
Db   3660 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3719

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
Db   3720 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3779

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
Db   3780 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3839

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
Db   3840 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3899

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
Db   3900 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3959

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
Db   3960 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4000


RESULT 7
US-09-740-211-13
; Sequence 13, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
```

```
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-13

  Query Match            100.0%;  Score 1001;  DB 10;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 4.3e-299;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3000 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 3059

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3060 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 3119

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3120 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 3179

Qy     181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3180 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 3239

Qy     241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3240 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 3299

Qy     301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3300 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 3359

Qy     361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3360 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 3419

Qy     421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3420 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 3479

Qy     481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3480 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 3539

Qy     541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3540 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 3599

Qy     601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3600 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 3659

Qy     661 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db    3660 ACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 3719

Qy     721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3720 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 3779

Qy     781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3780 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 3839

Qy     841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3840 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3899

Qy     901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3900 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3959

Qy     961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
          |||||||||||||||||||||||||||||||||||||||||||
Db    3960 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 4000


RESULT 8
US-10-095-718-3
; Sequence 3, Application US/10095718
; Patent No. US20020131956A1
; GENERAL INFORMATION:
;  APPLICANT: Walsh, Christopher
;  APPLICANT:  Chao, Hengjun
;  APPLICANT:  Burstein, Haim
;  APPLICANT:  Lynch, Carmel
;  APPLICANT:  Stepan, Tony
;  APPLICANT:  Munson, Keith
;  TITLE OF INVENTION: Adeno-Associated Virus Vectors Encoding Factor VIII and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 35052/204375
;  CURRENT APPLICATION NUMBER: US/10/095,718
;  CURRENT FILING DATE:  2002-03-12
;  PRIOR APPLICATION NUMBER: 09/689,430
;  PRIOR FILING DATE: 2001-08-22
;  PRIOR APPLICATION NUMBER: 60/158,780
;  PRIOR FILING DATE: 1999-10-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3
;   LENGTH: 7914
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: rAAV vector with canine B-domain deleted factor
;   OTHER INFORMATION: VIII
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (435)...(4730)
US-10-095-718-3

  Query Match            85.6%;  Score 857;  DB 12;  Length 7914;
  Best Local Similarity   91.0%;  Pred. No. 1.3e-254;
  Matches  911;  Conservative    0;  Mismatches   90;  Indels    0;  Gaps    0;

Qy       1 GCCCTTATACCGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGC 60
          |||||||||||||||||||| |||||||| ||||||||| ||||||||||||||||||||
Db    2966 GCCCTTATACCGTGGAGAACTGAATGAACACTTGGGACTCTTGGGGCCATATATAAGAGC 3025

Qy      61 AGAAGTTGAAGATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTT 120
          |||||||||||| |||||| |||||||||||| ||| ||||||||||||||| |||||
Db    3026 AGAAGTTGAAGACAATATCGTGGTAACTTTCAAAAACCAGGCCTCTCGTCCCTACTCCTT 3085

Qy     121 CTATTCTAGCCTTATTTCTTATGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAA 180
          |||||||| ||||||||||||| |||||| || | |||||||||||||||||| |||
Db    3086 CTATTCTAGTCTTATTTCTTATGACGAAGATGAGGGACAAGGAGCAGAACCTAGAAGAAA 3145
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Qy    181 CTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACC 240
Db   3146 GTTTGTCAACCCTAATGAAACCAAAATTTACTTTTGGAAAGTGCAGCATCATATGGCACC 3205

Qy    241 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAA 300
Db   3206 CACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTTTCTGATGTTGATTTGGAGAA 3265

Qy    301 AGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCC 360
Db   3266 AGATGTGCACTCAGGCTTGATTGGACCCCTTCTGATCTGCCGCAGTAACACACTGAACCC 3325

Qy    361 TGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGA 420
Db   3326 TGCTCATGGGAGACAAGTGACAGTGCAGGAGTTTGCCCTGGTTTTCACTATATTCGATGA 3385

Qy    421 GACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATAT 480
Db   3386 GACTAAGAGCTGGTACTTCACTGAAAACCTGGAAAGGAACTGTAGAGCTCCCTGCAATGT 3445

Qy    481 CCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACAT 540
Db   3446 CCAGAAGGAGGACCCTACTCTAAAAGAAAACTTCCGCTTCCATGCAATCAACGGCTATGT 3505

Qy    541 AATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCT 600
Db   3506 GAAGGATACACTCCCTGGCTTAGTAATGGCTCAGGATCAAAAGGTTCGATGGTATCTGCT 3565

Qy    601 CAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGT 660
Db   3566 CAGCATGGGCAGCAACGAAAACATTCATTCCATTCACTTCAGTGGACATGTGTTCACTGT 3625

Qy    661 ACGAAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGAC 720
Db   3626 ACGGAAAAAAAGAGGAATATAAAATGGCAGTCTACAACCTCTATCCAGGTGTTTTTGAGAC 3685

Qy    721 AGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCA 780
Db   3686 TGTGGAAATGCTACCATCCCAAGTTGGAATCTGGCGGATAGAATGCCTTATCGGCGAGCA 3745

Qy    781 TCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAGTGTCAGACTCCCCT 840
Db   3746 CCTGCAAGCCGGGATGAGCACTCTGTTTCTGGTGTACAGCAAGAAGTGTCAGACTCCACT 3805

Qy    841 GGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 900
Db   3806 GGGGATGGCTTCCGGACACATTAGAGATTTTCAGATTACAGCTTCAGGACAATATGGACA 3865

Qy    901 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 960
Db   3866 GTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAATGCCTGGAGCACCAA 3925

Qy    961 GGAGCCCTTTTCTTGGATCAAGGTGGATCTGTTGGCACCAA 1001
Db   3926 GGATCCCTTTTCCTGGATCAAGGTGGATCTCTTGGCACCGA 3966
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RESULT 9
US-09-880-107-2275
; Sequence 2275, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
; APPLICANT: Vockley, Joseph G.
; APPLICANT: Scherf, Uwe
; APPLICANT: Gene Logic, Inc.
; TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
; FILE REFERENCE: 44921-5028-WO
; CURRENT APPLICATION NUMBER: US/09/880,107
; CURRENT FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/211,379
; PRIOR FILING DATE: 2000-06-14
; PRIOR APPLICATION NUMBER: US 60/237,054
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 3950
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2275
; LENGTH: 6909
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20020142981A1 M16967
US-09-880-107-2275

Query Match 14.7%; Score 147.6; DB 10; Length 6909;
Best Local Similarity 49.5%; Pred. No. 4.5e-35;
Matches 479; Conservative 0; Mismatches 444; Indels 45; Gaps 2;

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Qy     11 CGTGGAGAACTAAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGCAGAAGTTGAA 70
Db   4990 CGAGGGGAGTATGAAGAGCATCTCGGAATTCTTGGTCCTATTATCAGAGCTGAAGTGGAT 5049

Qy     71 GATAATATCATGGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTTCTATTCTAGC 130
Db   5050 GATGTTATCCAAGTTCGTTTTAAAAATTTAGCATCCAGACCGTATTCTCTACATGCCCAT 5109

Qy    131 CTTATTTCTTATGAG----------GAAGATCAGAGGCAAGGAGCA 166
Db   5110 GGACTTTCCTATGAAAAATCATCAGAGGGAAAGACTTATGAAGATGACTCTCCTGAATGG 5169

Qy    167 GAACCTAGAAAAAACTTTGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAA 226
Db   5170 TTTAAGGAAGATAATGCTGTTCAGCCAAATAGCAGTTATACCTACGTATGGCATGCCACT 5229

Qy    227 CATCATATGGCACCCACTAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGAT 286
Db   5230 GAGCGATCAGGGCCAGAAAGTCCTGGCTCTGCCTGTCGGGCTTGGGCCTACTACTCAGCT 5289

Qy    287 GTTGACCTGGAAAAAGATGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACT 346
Db   5290 GTGAACCCAGAAAAAGATATTCACTCAGGCTTGATAGGTCCCCTCCTAATCTGCCAAAAA 5349

Qy    347 AACACACTGAACCCTGCTCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTC 406
Db   5350 GGAATACTACATAAGGACAGCAACATGCCTGTGGACATGAGAGAATTTGTCTTACTATTT 5409

Qy    407 ACCATCTTTGATGAGACCAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGG 466
Db   5410 ATGACCTTTGATGAAAAGAAGAGCTGGTACTATGAAAAGAAGTCCCGAAGTTCTTGGAGA 5469

Qy    467 GCTCCCTGCAATATCCAGATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCA 526
Db   5470 CTCACATCCTCAGAAATGAAAAAATCCCATGAGTTTCACGCCATTAATGGG--------- 5520

Qy    527 ATCAATGGCTACATAATGGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATT 586
Db   5521 ----------ATGATCTACAGCTTGCCTGGCCTGAAAATGTATGAGCAAGAGTGGGTG 5568

Qy    587 CGATGGTATCTGCTCAGCATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGA 646
Db   5569 AGGTTACACCTGCTGAACATAGGCGGCTCCCAAGACATTCACGTGGTTCACTTTCACGGC 5628

Qy    647 CATGTGTTCACTGTACGAAAAAAAGAGGAGTATAAAATGGCACTGTACAATCTCTATCCA 706
Db   5629 CAGACCTTGCTGGAAAATGGCAATAAACAGCACCAGTTAGGGGTCTGGCCCCTTCTGCCT 5688

Qy    707 GGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAATGC 766
Db   5689 GGTTCATTTAAAACTCTTGAAATGAAGGCATCAAAACCTGGCTGGTGGCTCCTAAACACA 5748

Qy    767 CTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAATAAG 826
Db   5749 GAGGTTGGAGAAAACCAGAGAGCAGGGATGCAAACGCCATTTCTTATCATGGACAGAGAC 5808

Qy    827 TGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCTTCA 886
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Db   5809 TGTAGGATGCCAATGGGACTAAGCACTGGTATCATATCTGATTCACAGATCAAGGCTTCA 5868

Qy    887 GGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATCAAT 946
          |        || | |||  || |   | || ||| |  |  ||   |||||    |||
Db   5869 GAGTTTCTGGGTTACTGGGAGCCCAGATTAGCAAGATTAAACAATGGTGGATCTTATAAT 5928

Qy    947 GCCTGGAG 954
          || |||||
Db   5929 GCTTGGAG 5936


RESULT 10
US-09-970-966-175
; Sequence 175, Application US/09970966
; Patent No. US20020173638A1
; GENERAL INFORMATION:
;  APPLICANT: Stolk, John A.
;  APPLICANT:  Molesh, David Alan
;  APPLICANT:  Fling, Steven P.
;  APPLICANT:  Xu, Jiangchun
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.484C6
;  CURRENT APPLICATION NUMBER: US/09/970,966
;  CURRENT FILING DATE:  2001-10-02
;  NUMBER OF SEQ ID NOS: 215
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 175
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-970-966-175

  Query Match             13.5%;  Score 135;  DB 9;  Length 3321;
  Best Local Similarity   50.7%;  Pred. No. 2.3e-31;
  Matches  398;  Conservative    0;  Mismatches  360;  Indels   27;  Gaps    2;

Qy      22 AAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGCAGAAGTTGAAGATAATATCAT 81
           | | |||||| |||||  | || || ||| |  |    ||||| |||| ||| ||  |||
Db    2403 AGAAGAACATCTGGGAATTCTAGGTCCACAACTTCATGCAGATGTTGGAGACAAAGTCAA 2462

Qy      82 GGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTTCTATTCTAGCCTTATTTCTTA 141
             | |  || | |||   |||| |  | ||||| ||  |  || |
Db    2463 AATTATCTTTAAAAACATGGCCACAAGGCCCTACTCAATACATGCCC------------- 2509

Qy     142 TGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAACTTTGTCAAGCCTAATGAAAC 201
                ||  |   |||  ||      ||| |   |           ||   ||||||
Db    2510 --------ATGGGGTACAAACAGAGAGTTCTACAGTTACTCCAACATTACCAGGTGAAAC 2561

Qy     202 CAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACCCACTAAAGATGAGTTTGACTG 261
           |||||| | |||||| | | | |       | | |    | ||| || | ||  ||
Db    2562 TCTCACTTACGTATGGAAAATCCCAGAAAGATCTGGAGCTGGAACAGAGGATTCTGCTTG 2621

Qy     262 CAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAAAGATGTGCACTCAGGCCTGAT 321
            |   | ||||||||||  ||   ||| || |   |  || ||  |  ||   ||  | ||
Db    2622 TATTCCATGGGCTTATTATTCAACTGTGGATCAAGTTAAGGACCTCTACAGTGGATTAAT 2681

Qy     322 TGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCCTGCTCATGGGAGACAAGTGAC 381
           ||| |||||  | || || |  |          ||||        ||  |||     |
Db    2682 TGGCCCCCTGATTGTTTGTCGAAGACCTTACTTGAAAGTATTCAATCCCAGAAGGAAG-- 2739

Qy     382 AGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGAGACCAAAAGCTGGTACTTCAC 441
           | |||||||||  ||  | ||     | ||||||||||  ||  |||||||
Db    2740 ----CTGGAATTTGCCCTTCTGTTTCTAGTTTTTGATGAGAATGAATCTTGGTACTTAGA 2795

Qy     442 TGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATATCCAGATGGAAGATCCCACTTT 501
           ||| || ||  ||| | |||        |||  ||  |  | |  || |||      ||
Db    2796 TGACAACATCAAAACATACTCTGATCACCCCGAGAAAGTAAACAAAGATGATGAGGAATT 2855

Qy     502 TAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACATAATGGATACACTACCTGGCTT 561
           | ||| |   ||    | ||||| || |||||    ||  | |  |  ||||  ||| |
Db    2856 CATAGAAAGCAATAAAATGCATGCTATTAATGGAAGAATGTTTGGAAACCTACAAGGCCT 2915
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Qy     562 AGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAA 621
             ||||     ||   |     |    |||||||||  |  | |||||||   |  | |
Db    2916 CACAATGCACGTGGGAGATGAAGTCAACTGGTATCTGATGGGAATGGGCAATGAAATAGA 2975

Qy     622 CATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAA 681
           | | ||  || | |||||    || |||   |||           | |  |  |  ||||
Db    2976 CTTACACACTGTACATTTTCACGGCCATAGCTTCCAATACAAGCACAGGGGAGTTTATAG 3035

Qy     682 AATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAA 741
              |  | |   |  | |  || ||    |   | ||  | |||||||| |||   |
Db    3036 TTCTGATGTCTTTGACATTTTCCCTGGAACATACCAAACCCTAGAAATGTTTCCAAGAAC 3095

Qy     742 AGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCAC 801
           | ||||||||||     |  | |||| | |   ||| ||  | |||||||| |||   ||
Db    3096 ACCTGGAATTTGGTTACTCCACTGCCATGTGACCGACCACATTCATGCTGGAATGGAAAC 3155

Qy     802 ACTTT 806
           ||
Db    3156 CACTT 3160


RESULT 11
US-09-825-294-175
; Sequence 175, Application US/09825294
; Patent No. US20020004491A1
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Stolk, John A.
;  APPLICANT:  Algate, Paul A.
;  APPLICANT:  Fling, Steven P.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
;  TITLE OF INVENTION:  THERAPY AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.484C5
;  CURRENT APPLICATION NUMBER: US/09/825,294
;  CURRENT FILING DATE:  2001-04-03
;  NUMBER OF SEQ ID NOS: 215
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 175
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-825-294-175

  Query Match             13.5%;  Score 135;  DB 10;  Length 3321;
  Best Local Similarity   50.7%;  Pred. No. 2.3e-31;
  Matches  398;  Conservative    0;  Mismatches  360;  Indels   27;  Gaps    2;

Qy      22 AAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGCAGAAGTTGAAGATAATATCAT 81
           | | |||||| |||||  | || || ||| |  |    ||||| |||| ||| ||  |||
Db    2403 AGAAGAACATCTGGGAATTCTAGGTCCACAACTTCATGCAGATGTTGGAGACAAAGTCAA 2462

Qy      82 GGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTTCTATTCTAGCCTTATTTCTTA 141
             | |  || | |||   |||| |  | ||||| ||  |  || |
Db    2463 AATTATCTTTAAAAACATGGCCACAAGGCCCTACTCAATACATGCCC------------- 2509

Qy     142 TGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAACTTTGTCAAGCCTAATGAAAC 201
                ||  |   |||  ||      ||| |   |           ||   ||||||
Db    2510 --------ATGGGGTACAAACAGAGAGTTCTACAGTTACTCCAACATTACCAGGTGAAAC 2561

Qy     202 CAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACCCACTAAAGATGAGTTTGACTG 261
           |||||| | |||||| | | | |       | | |    | ||| || | ||  ||
Db    2562 TCTCACTTACGTATGGAAAATCCCAGAAAGATCTGGAGCTGGAACAGAGGATTCTGCTTG 2621

Qy     262 CAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAAAGATGTGCACTCAGGCCTGAT 321
            |   | ||||||||||  ||   ||| || |   |  || ||  |  ||   ||  | ||
Db    2622 TATTCCATGGGCTTATTATTCAACTGTGGATCAAGTTAAGGACCTCTACAGTGGATTAAT 2681

Qy     322 TGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCCTGCTCATGGGAGACAAGTGAC 381
           ||| |||||  | || || |  |          ||||        ||  |||     |
Db    2682 TGGCCCCCTGATTGTTTGTCGAAGACCTTACTTGAAAGTATTCAATCCCAGAAGGAAG-- 2739

Qy     382 AGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGAGACCAAAAGCTGGTACTTCAC 441
           | |||||||||  ||  | ||     | ||||||||||  ||  |||||||
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Db   2740 ----CTGGAATTTGCCCTTCTGTTTCTAGTTTTTGATGAGAATGAATCTTGGTACTTAGA 2795

Qy    442 TGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATATCCAGATGGAAGATCCCACTTT 501
          ||| || ||  ||| | |||         |||   ||  |  | |  || ||| |     ||
Db   2796 TGACAACATCAAAACATACTCTGATCACCCCGAGAAAGTAAACAAAGATGATGAGGAATT 2855

Qy    502 TAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACATAATGGATACACTACCTGGCTT 561
          | ||| |   ||    | ||||| || |||||    ||  | |  |  ||||  ||| |
Db   2856 CATAGAAAGCAATAAAATGCATGCTATTAATGGAAGAATGTTTGGAAACCTACAAGGCCT 2915

Qy    562 AGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAA 621
            ||||     ||   |      |    |||||||||| |  | |||||||   |   | |
Db   2916 CACAATGCACGTGGGAGATGAAGTCAACTGGTATCTGATGGGAATGGGCAATGAAATAGA 2975

Qy    622 CATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAA 681
          | | ||  || | |||||    || |||   |||          | |  |  |  ||||
Db   2976 CTTACACACTGTACATTTTCACGGCCATAGCTTCCAATACAAGCACAGGGGAGTTTATAG 3035

Qy    682 AATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAA 741
            |   | |    |  | |  || ||    |   | ||  | |||||||| |||   |
Db   3036 TTCTGATGTCTTTGACATTTTCCCTGGAACATACCAAACCCTAGAAATGTTTCCAAGAAC 3095

Qy    742 AGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCAC 801
          | ||||||||||||    |  | |||| | |   ||| ||  | |||||||| |||   ||
Db   3096 ACCTGGAATTTGGTTACTCCACTGCCATGTGACCGACCACATTCATGCTGGAATGGAAAC 3155

Qy    802 ACTTT 806
            ||
Db   3156 CACTT 3160


RESULT 12
US-09-880-107-2253
; Sequence 2253, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,107
;  CURRENT FILING DATE:  2001-06-14
;  PRIOR APPLICATION NUMBER: US 60/211,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2253
;   LENGTH: 3321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020142981A1 M13699
US-09-880-107-2253

  Query Match             13.5%;  Score 135;  DB 10;  Length 3321;
  Best Local Similarity   50.7%;  Pred. No. 2.3e-31;
  Matches  398;  Conservative    0;  Mismatches  360;  Indels   27;  Gaps    2;

Qy     22 AAATGAACATTTGGGACTCCTGGGGCCATATATAAGAGCAGAAGTTGAAGATAATATCAT 81
          | | |||||| ||||| | || || ||| |  |     ||||| |||| ||| ||  |||
Db   2403 AGAAGAACATCTGGGAATTCTAGGTCCACAACTTCATGCAGATGTTGGAGACAAAGTCAA 2462

Qy     82 GGTAACTTTCAGAAATCAGGCCTCTCGTCCCTATTCCTTCTATTCTAGCCTTATTTCTTA 141
            | |  || | |||   |||| |  | ||||| ||  |  || |
Db   2463 AATTATCTTTAAAAACATGGCCACAAGGCCCTACTCAATACATGCCC------------- 2509

Qy    142 TGAGGAAGATCAGAGGCAAGGAGCAGAACCTAGAAAAAACTTTGTCAAGCCTAATGAAAC 201
                ||  |   |||  ||      ||| |   |           ||   ||||||
Db   2510 --------ATGGGGTACAAACAGAGAGTTCTACAGTTACTCCAACATTACCAGGTGAAAC 2561

Qy    202 CAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACCCACTAAAGATGAGTTTGACTG 261
          ||||||  | ||||||  | | | |        | | |     | ||| || | ||  ||
Db   2562 TCTCACTTACGTATGGAAAATCCCAGAAAGATCTGGAGCTGGAACAGAGGATTCTGCTTG 2621

Qy    262 CAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAAAGATGTGCACTCAGGCCTGAT 321
            |   | ||||||||||  ||   ||| || |  |  || ||  |  ||   || | ||
Db   2622 TATTCCATGGGCTTATTATTCAACTGTGGATCAAGTTAAGGACCTCTACAGTGGATTAAT 2681

Qy    322 TGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCCTGCTCATGGGAGACAAGTGAC 381
          ||| |||||  | || || |  |          ||||         ||   |||     |
Db   2682 TGGCCCCCTGATTGTTTGTCGAAGACCTTACTTGAAAGTATTCAATCCCAGAAGGAAG-- 2739

Qy    382 AGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGAGACCAAAAGCTGGTACTTCAC 441
              | ||||||||| ||  | ||      | ||||||||||   ||   ||||||||
Db   2740 ----CTGGAATTTGCCCTTCTGTTTCTAGTTTTTGATGAGAATGAATCTTGGTACTTAGA 2795

Qy    442 TGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATATCCAGATGGAAGATCCCACTTT 501
          |||| || ||  ||| | |||         |||   ||  |  | |  || ||| |     ||
Db   2796 TGACAACATCAAAACATACTCTGATCACCCCGAGAAAGTAAACAAAGATGATGAGGAATT 2855

Qy    502 TAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACATAATGGATACACTACCTGGCTT 561
          | ||| |   ||    | ||||| || |||||    ||  | |  |  ||||  ||| |
Db   2856 CATAGAAAGCAATAAAATGCATGCTATTAATGGAAGAATGTTTGGAAACCTACAAGGCCT 2915

Qy    562 AGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCTCAGCATGGGCAGCAATGAAAA 621
            ||||     ||   |      |    |||||||||| |  | |||||||   |   | |
Db   2916 CACAATGCACGTGGGAGATGAAGTCAACTGGTATCTGATGGGAATGGGCAATGAAATAGA 2975

Qy    622 CATCCATTCTATTCATTTCAGTGGACATGTGTTCACTGTACGAAAAAAAGAGGAGTATAA 681
          | | ||  || | |||||    || |||   |||          | |  |  |  ||||
Db   2976 CTTACACACTGTACATTTTCACGGCCATAGCTTCCAATACAAGCACAGGGGAGTTTATAG 3035

Qy    682 AATGGCACTGTACAATCTCTATCCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAA 741
            |   | |    |  | |  || ||    |   | ||  | |||||||| |||   |
Db   3036 TTCTGATGTCTTTGACATTTTCCCTGGAACATACCAAACCCTAGAAATGTTTCCAAGAAC 3095

Qy    742 AGCTGGAATTTGGCGGGTGGAATGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCAC 801
          | ||||||||||||    |  | |||| | |   ||| ||  | |||||||| |||   ||
Db   3096 ACCTGGAATTTGGTTACTCCACTGCCATGTGACCGACCACATTCATGCTGGAATGGAAAC 3155

Qy    802 ACTTT 806
            ||
Db   3156 CACTT 3160


RESULT 13
US-09-917-800A-1539
; Sequence 1539, Application US/09917800A
; Patent No. US20020119462A1
; GENERAL INFORMATION:
;  APPLICANT: Mendrick, Donna
;  APPLICANT:  Porter, Mark
;  APPLICANT:  Johnson, Kory
;  APPLICANT:  Castle, Arthur
;  APPLICANT:  Elashoff, Michael
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Molecular Toxicology Modeling
;  FILE REFERENCE: 44921-5038-US
;  CURRENT APPLICATION NUMBER: US/09/917,800A
;  CURRENT FILING DATE:  2001-07-31
;  PRIOR APPLICATION NUMBER: US 60/222,040
;  PRIOR FILING DATE: 2000-07-31
;  PRIOR APPLICATION NUMBER: US 60/222,880
;  PRIOR FILING DATE: 2000-11-02
;  PRIOR APPLICATION NUMBER: US 60/290,029
;  PRIOR FILING DATE: 2001-05-11
;  PRIOR APPLICATION NUMBER: US 60/290,645
;  PRIOR FILING DATE: 2001-05-15
;  PRIOR APPLICATION NUMBER: US 60/292,336
;  PRIOR FILING DATE: 2001-05-22
;  PRIOR APPLICATION NUMBER: US 60/295,798
;  PRIOR FILING DATE: 2001-06-06
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;  PRIOR APPLICATION NUMBER: US 60/297,457
;  PRIOR FILING DATE: 2001-06-13
;  PRIOR APPLICATION NUMBER: US 60/298,884
;  PRIOR FILING DATE: 2001-06-19
;  PRIOR APPLICATION NUMBER: US 60/303,459
;  PRIOR FILING DATE: 2001-07-09
;  NUMBER OF SEQ ID NOS: 1740
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1539
;   LENGTH: 3700
;   TYPE: DNA
;   ORGANISM: Rattus norvegicus
;   FEATURE:
;   OTHER INFORMATION: Genbank Accession No. US20020119462A1 NM_012532
US-09-917-800A-1539

  Query Match             10.6%;  Score 106;  DB 10;  Length 3700;
  Best Local Similarity   51.7%;  Pred. No. 2.3e-22;
  Matches  241;  Conservative    0;  Mismatches  225;  Indels    0;  Gaps    0;

Qy     184 TGTCAAGCCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACCCAC 243
           |||    || || ||||||   || ||    ||||  || |   |  | |||| ||||||
Db    1521 TGTGGCTCCCAAAGAAACCTTTACGTATGAATGGACTGTCCCCAAAGAAATGGGACCCAC 1580

Qy     244 TAAAGATGAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAAAGA 303
           | | |  ||   ||  |||  | |   |  |||| |||| ||||||||    |||||
Db    1581 TTATGCAGATCCTGTGTGCCTATCTAAGATGTATTATTCTGGAGTTGACCTCACCAAAGA 1640

Qy     304 TGTGCACTCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCCTGC 363
           | |     | || || ||||| ||  |     | ||| | |    ||   |     | |
Db    1641 TATATTTACTGGGCTTATTGGGCCAATGAAAATATGCAAGAAAGGCAGCTTACTTGCAGA 1700

Qy     364 TCATGGGAGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGAGAC 423
           |   |     |||    ||   |||| ||     |||||   ||  | ||||||||||
Db    1701 TGGGAGACAGAAAGATGTAGACAAGGAGTTCTACTTGTTTGCAACAGTGTTTGATGAGAA 1760

Qy     424 CAAAAGCTGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATATCCA 483
             || |    |||   ||| ||||||   ||    |   |  || ||    ||| |  |
Db    1761 TGAGAGTTTACTCTTGGATGATAATATCAGAATGTTCACAACTGCACCTGAGAATGTGGA 1820

Qy     484 GATGGAAGATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACATAAT 543
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Db    1821 CAAGGAAGATGAAGACTTTCAGGAGTCCAACAAGATGCACTCCATGAATGGATTCATGTA 1880

Qy     544 GGATACACTACCTGGCTTAGTAATGGCTCAGGATCAAAGGATTCGATGGTATCTGCTCAG 603
             |  |  || |||||| |    |||   |  |   ||   ||    |||||| || ||||
Db    1881 TGGCAATCTGCCTGGCCTCAATATGTGCCTAGGAGAATCCATCGTGTGGTATTTGTTCAG 1940

Qy     604 CATGGGCAGCAATGAAAACATCCATTCTATTCATTTCAGTGGACAT 649
           |   || |   | | | || | |||   ||  | ||    ||| ||
Db    1941 CGCTGGAAATGAGGCAGACGTGCATGGGATATACTTTTCAGGAAAT 1986


RESULT 14
US-09-960-352-13850
; Sequence 13850, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 13850
;   LENGTH: 357
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   OTHER INFORMATION: Clone ID: 59-LIB34-008-Q1-E1-G12
US-09-960-352-13850

  Query Match             6.9%;  Score 69.2;  DB 10;  Length 357;
  Best Local Similarity   54.8%;  Pred. No. 1.4e-11;
  Matches  137;  Conservative    0;  Mismatches  113;  Indels    0;  Gaps    0;

Qy     704 CCAGGTGTTTTTGAGACAGTGGAAATGTTACCATCCAAAGCTGGAATTTGGCGGGTGGAA 763
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Db      34 CCTGGTTCATTTAAAACTCTTGAAATGAAGGCATCAAAACCTGGCTGGTGGCTCCTAGAC 93

Qy     764 TGCCTTATTGGCGAGCATCTACATGCTGGGATGAGCACACTTTTTCTGGTGTACAGCAAT 823
               |||| ||   ||     || ||||||   ||||  |||||  |      ||
Db      94 ACGGAAGTTGGAGAAATTCAGAGAGCAGGGATGCAGACACCATTTCTCATTGTAGACAGA 153

Qy     824 AAGTGTCAGACTCCCCTGGGAATGGCTTCTGGACACATTAGAGATTTTCAGATTACAGCT 883
            | ||| |||  ||  ||||| |     |||| |  ||    || |  |||||    |||
Db     154 GAATGTAAGATGCCAATGGGACTAAGCACTGGCCTGATAGCTGACTCACAGATCCAGGCT 213

Qy     884 TCAGGACAATATGGACAGTGGGCCCCAAAGCTGGCCAGACTTCATTATTCCGGATCAATC 943
           || |     |  ||  | ||||  || ||  | || ||  |  |  ||   ||||||  |
Db     214 TCTGAGTTTTGGGGTTATTGGGAACCCAAATTAGCAAGGTTAAACAATGGTGGATCATAC 273

Qy     944 AATGCCTGGA 953
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Db     274 AATGCTTGGA 283


RESULT 15
US-09-960-352-12959
; Sequence 12959, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 12959
;   LENGTH: 389
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   FEATURE:
;   NAME/KEY: unsure
;   LOCATION: (360)
;   OTHER INFORMATION: unsure at all n locations
;   OTHER INFORMATION: Clone ID: 55-LIB34-039-Q1-E1-F12
US-09-960-352-12959

  Query Match             6.5%;  Score 64.8;  DB 10;  Length 389;
  Best Local Similarity   51.0%;  Pred. No. 3.5e-10;
  Matches  181;  Conservative    0;  Mismatches  168;  Indels    6;  Gaps    1;

Qy     191 CCTAATGAAACCAAAACTTACTTTTGGAAAGTGCAACATCATATGGCACCCACTAAAGAT 250
           ||   ||||||    |||||| | |||||| | | | |       | | |    |  |||
Db      41 CCAGGTGAAACTCGCACTTACATATGGAAAATCCCAGAAAGATCTGGAGCTGGAAGGGAT 100

Qy     251 GAGTTTGACTGCAAAGCCTGGGCTTATTTCTCTGATGTTGACCTGGAAAAAGATGTGCAC 310
           || | |   || |   | |||| ||| |  ||   ||| || | ||  || ||| |
Db     101 GATTCTCCTTGTATTCCATGGGTTTACTATTCAACTGTGGATCGGGTTAAGGATCTCTTT 160

Qy     311 TCAGGCCTGATTGGACCCCTTCTGGTCTGCCACACTAACACACTGAACCCTGCTCATGGG 370
             ||  | ||||| || ||  | || || |  |   ||    ||||        ||
Db     161 AGTGGATTAATTGGCCCACTGATTGTTTGTCGGAAGCACTATTTGAAAGTATTCAATCCC 220

Qy     371 AGACAAGTGACAGTACAGGAATTTGCTCTGTTTTTCACCATCTTTGATGAGACCAAAAGC 430
           | | |         || ||||||| | ||  | ||     | ||||||||||   ||
Db     221 ATAAAGAA------ACTGGAATTTTCCCTTCTGTTTCTTGTTTTTGATGAGAATGAATCT 274
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Qy    431 TGGTACTTCACTGAAAATATGGAAAGAAACTGCAGGGCTCCCTGCAATATCCAGATGGAA 490
          ||||||||   ||| |||||  ||| | |||         ||    ||  |  | |  |
Db    275 TGGTACTTAGATGACAATATCAAAACATACTCTGATCACCCTGAGAAAGTAGACAAAGCC 334

Qy    491 GATCCCACTTTTAAAGAGAATTATCGCTTCCATGCAATCAATGGCTACATAATGG 545
           ||      || |  || |   ||    | ||||| || |||||  | ||  | |
Db    335 AATGAGGAATTCATGGAAAGCAATANAATGCATGCTATTAATGGAAAAATGTTTG 389
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GenCore version 5.1.3

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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.		Score	% Query Match	Length	DB	ID	Description
1		1001	100.0	11933	4	US-09-470-618-13	Sequence 13, Appl
2		1001	100.0	11933	4	US-09-364-862-13	Sequence 13, Appl
3	c	926.6	92.6	46819	4	US-09-453-702B-72	Sequence 72, Appl
4	c	575.2	57.5	61663	4	US-09-453-702B-62	Sequence 62, Appl
5		309	30.9	309	3	US-09-023-221A-6	Sequence 6, Appli
6		309	30.9	309	4	US-09-282-352A-6	Sequence 6, Appli
7		298.8	29.9	760	3	US-09-023-221A-1	Sequence 1, Appli
8		298.8	29.9	760	4	US-09-282-352A-1	Sequence 1, Appli
9		256.4	25.6	378	3	US-09-023-221A-21	Sequence 21, Appl
10		256.4	25.6	378	4	US-09-282-352A-21	Sequence 21, Appl
11		252	25.2	309	3	US-09-023-221A-22	Sequence 22, Appl
12		252	25.2	309	4	US-09-282-352A-22	Sequence 22, Appl
13		224.2	22.4	309	3	US-09-023-221A-5	Sequence 5, Appli
14		224.2	22.4	309	4	US-09-282-352A-5	Sequence 5, Appli
15	c	59.2	5.9	38584	4	US-09-453-702B-50	Sequence 50, Appl
16	c	55.8	5.6	26173	4	US-09-453-702B-69	Sequence 69, Appl
17		55.8	5.6	48908	4	US-09-453-702B-137	Sequence 137, App
18		55.2	5.5	45175	4	US-09-453-702B-116	Sequence 116, App
19	c	54	5.4	38155	4	US-09-453-702B-79	Sequence 79, Appl
20		39	3.9	9048	3	US-08-973-273-4	Sequence 4, Appli
21		37.2	3.7	595	4	US-09-385-982-25	Sequence 25, Appl
22		35.6	3.6	22306	4	US-09-453-702B-251	Sequence 251, App
23		35.6	3.6	34063	4	US-09-453-702B-96	Sequence 96, Appl
24		35.4	3.5	49795	4	US-09-453-702B-60	Sequence 60, Appl
25		35	3.5	1727	4	US-09-071-035-295	Sequence 295, App
26		35	3.5	1839	4	US-09-071-035-293	Sequence 293, App
27		34.4	3.4	112132	4	US-09-741-150-3	Sequence 3, Appli
28	c	34.2	3.4	16950	4	US-09-453-702B-166	Sequence 166, App
29		34	3.4	3680	4	US-09-647-390-15	Sequence 15, Appl
30		33.6	3.4	1611	6	5213972-6	Patent No. 5213972
31		33.6	3.4	1817	2	US-08-743-637B-1	Sequence 1, Appli
32		33.6	3.4	1817	3	US-08-526-840B-1	Sequence 1, Appli
33	c	33.4	3.3	1391	2	US-08-950-168-2	Sequence 2, Appli
34	c	33.4	3.3	1391	4	US-09-365-705-2	Sequence 2, Appli
35	c	33.4	3.3	1404	4	US-09-257-179-34	Sequence 34, Appl
36		33.4	3.3	9711	4	US-08-961-527-167	Sequence 167, App
37		33	3.3	566	4	US-09-328-111-452	Sequence 452, App
38		33	3.3	2400	4	US-08-963-901-1	Sequence 1, Appli
39		33	3.3	2400	4	US-08-963-901-5	Sequence 5, Appli
40	c	33	3.3	2796	1	US-08-261-677-8	Sequence 8, Appli
41	c	33	3.3	2796	1	US-08-384-556A-4	Sequence 4, Appli
42	c	33	3.3	2796	2	US-08-331-355A-8	Sequence 8, Appli
43	c	33	3.3	2796	3	US-08-777-147-5	Sequence 5, Appli
44	c	33	3.3	2796	3	US-09-157-077-8	Sequence 8, Appli
45	c	33	3.3	2796	5	PCT-US94-12364-8	Sequence 8, Appli

ALIGNMENTS

RESULT 1
US-09-470-618-13
; Sequence 13, Application US/09470618
; Patent No. 6200560
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/09/470,618
; CURRENT FILING DATE: 1999-12-22
; EARLIER APPLICATION NUMBER: 09/364,862
; EARLIER FILING DATE: 1999-07-30
; EARLIER APPLICATION NUMBER: 60/125,974
; EARLIER FILING DATE: 1999-03-24
; EARLIER APPLICATION NUMBER: 60/104,994
; EARLIER FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
; LENGTH: 11933
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-470-618-13

Query Match 100.0%; Score 1001; DB 4; Length 11933;
Best Local Similarity 100.0%; Pred. No. 2.1e-283;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8700 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 8759

Qy      61 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8760 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 8819

Qy     121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8820 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 8879

Qy     181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8880 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 8939

Qy     241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    8940 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 8999

Qy     301 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9000 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 9059

Qy     361 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9060 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 9119

Qy     421 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9120 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 9179

Qy     481 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9180 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 9239

Qy     541 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9240 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 9299

Qy     601 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9300 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 9359

Qy     661 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9360 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 9419

Qy     721 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9420 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 9479

Qy     781 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9480 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 9539

Qy     841 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9540 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 9599

Qy     901 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9600 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 9659

Qy     961 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
           |||||||||||||||||||||||||||||||||||||||||
Db    9660 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 9700


RESULT 2
US-09-364-862-13
; Sequence 13, Application US/09364862
; Patent No. 6221349
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT: Colosi, Peter C.
;  TITLE OF INVENTION: ADENO-ASSOCIATED VECTORS FOR EXPRESSION OF FACTOR VIII
;  TITLE OF INVENTION: BY TARGET
;  TITLE OF INVENTION: CELLS
;  FILE REFERENCE: AVIGEN-03743
;  CURRENT APPLICATION NUMBER: US/09/364,862
;  CURRENT FILING DATE: 1999-07-30
;  EARLIER APPLICATION NUMBER: 60/125,974
;  EARLIER FILING DATE: 1999-03-24
;  EARLIER APPLICATION NUMBER: 60/104,994
;  EARLIER FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 14
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  synthetic
US-09-364-862-13

  Query Match             100.0%;  Score 1001;  DB 4;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 2.1e-283;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8700 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 8759

Qy      61 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8760 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 8819

Qy     121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8820 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 8879

Qy     181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8880 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 8939

Qy     241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8940 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 8999

Qy     301 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9000 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 9059

Qy     361 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9060 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 9119

Qy     421 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9120 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 9179

Qy     481 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9180 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 9239

Qy     541 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9240 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 9299

Qy     601 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9300 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 9359

Qy     661 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9360 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 9419

Qy     721 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9420 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 9479

Qy     781 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9480 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 9539

Qy     841 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9540 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 9599

Qy     901 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9600 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 9659
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Qy    961 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          ||||||||||||||||||||||||||||||||||||||||||
Db   9660 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 9700


RESULT 3
US-09-453-702B-72/c
; Sequence 72, Application US/09453702B
; Patent No. 6365723
;   GENERAL INFORMATION:
;        APPLICANT: Blattner, Frederick R.
;                   Burland,     Valerie
;                   Perna,  Nicole T.
;                   Plunkett, Guy
;                   Welch,  Rod
;        TITLE OF INVENTION: No. 6365723el Sequences of E. coli O157
;        NUMBER OF SEQUENCES: 265
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Quarles & Brady
;             STREET: 1 South Pinckney Street
;             CITY: Madison
;             STATE: WI
;             COUNTRY: US
;             ZIP: 53701-2113
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: Word Perfect 8.0
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/453,702B
;             FILING DATE: 03-Dec-1999
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 60/110,955
;             FILING DATE: 04-DEC-1998
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Seay, Nicholas J.
;             REGISTRATION NUMBER: 27386
;             REFERENCE/DOCKET NUMBER: 960296.95017
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (608) 251-5000
;             TELEFAX: (608) 251-9166
;   INFORMATION FOR SEQ ID NO: 72:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 46819
;             TYPE: nucleic acid
;             STRANDEDNESS: double
;             TOPOLOGY: linear
;        MOLECULE TYPE: DNA  (genomic)
;        SEQUENCE DESCRIPTION: SEQ ID NO: 72:
US-09-453-702B-72

  Query Match             92.6%;  Score 926.6;  DB 4;  Length 46819;
  Best Local Similarity   97.2%;  Pred. No. 2.5e-261;
  Matches  975;  Conservative    0;  Mismatches   24;  Indels    4;  Gaps    3;

Qy      1 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 60
          || |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  10601 AATGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATTGTGTTAGAAAAC 10542

Qy     61 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 120
          |||||||| ||||||||||||||||||||||||||||| ||||||||||||||||||||
Db  10541 GATAACACTGTGTTAATAAAAGGACTTAAAAAGGTTGTGAATGTTAAATTCTCAAGAAAT 10482

Qy    121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  10481 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 10422

Qy    181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
          |||||||||||||||||||||||||||||||||||||||||||||||||||| ||||||
Db  10421 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCACGAACAT 10362

Qy    241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
          ||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db  10361 AAAAGATATTACTATGCCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 10302

Qy    301 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 360
          |||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||
Db  10301 ACACAAATCTTTCCACGCTAAATCATCACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 10242

Qy    361 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTT-TTAATTATTCCC 419
          |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db  10241 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTATTAATTATTCCC 10182

Qy    420 GGACACTCCCGCAGAGAAG-TTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATAC 478
          |||||||||||||||| || ||||| |||||||||| ||||||||||||||||||||||
Db  10181 GGACACTCCCGCAGAGGAGTTTCCCTGTCAGGGCTGCGGACATAGTTAATCCGGGAATAC 10122

Qy    479 AATGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACT 538
          | ||||||| | ||||| |||||||||||||||||||||||||||||| |||||||||
Db  10121 AGTGACGATCCTTCGCATCTGACATACATTAATAAATATTAACAATATGAGATTTCAACT 10062

Qy    539 CATTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATC 598
          |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db  10061 CATTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTTAAAAAGCATC 10002

Qy    599 GGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGA 658
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  10001 GGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGA 9942

Qy    659 TGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACC 718
          ||||| |||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db   9941 TGTGCCCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAACACCAAAGGAAACC 9882

Qy    719 ATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAA 778
          |||||||| |||||||||| ||||||||||||||||||||||||||||||||||| |||
Db   9881 ATCACCCACCATTTCTTCGTCTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCTAAA 9822

Qy    779 ATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGA 838
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9821 ATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGA 9762

Qy    839 TTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCA 898
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9761 TTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCA 9702

Qy    899 CAATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAAT 958
          ||||||||||||||| ||||||| |||||||||| ||||||||||||||||||||||||
Db   9701 CAATAATTGCATGAGCTGCCCAT--CGATATGGGCAGCTCTATCTGCACTGCTCATTAAT 9644

Qy    959 ATACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          |||||||||||||||||||||||||||||||||||||||||||
Db   9643 ATACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 9601


RESULT 4
US-09-453-702B-62/c
; Sequence 62, Application US/09453702B
; Patent No. 6365723
;   GENERAL INFORMATION:
;        APPLICANT: Blattner, Frederick R.
;                   Burland,     Valerie
;                   Perna,  Nicole T.
;                   Plunkett, Guy
;                   Welch,  Rod
;        TITLE OF INVENTION: No. 6365723el Sequences of E. coli O157
;        NUMBER OF SEQUENCES: 265
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Quarles & Brady
;             STREET: 1 South Pinckney Street
;             CITY: Madison
;             STATE: WI
;             COUNTRY: US
;             ZIP: 53701-2113
;        COMPUTER READABLE FORM:
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;             MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: Word Perfect 8.0
;       CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/453,702B
;             FILING DATE: 03-Dec-1999
;             CLASSIFICATION: <Unknown>
;       PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 60/110,955
;             FILING DATE: 04-DEC-1998
;       ATTORNEY/AGENT INFORMATION:
;             NAME: Seay, Nicholas J.
;             REGISTRATION NUMBER: 27386
;             REFERENCE/DOCKET NUMBER: 960296.95017
;       TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (608) 251-5000
;             TELEFAX: (608) 251-9166
;   INFORMATION FOR SEQ ID NO: 62:
;       SEQUENCE CHARACTERISTICS:
;             LENGTH: 61663
;             TYPE: nucleic acid
;             STRANDEDNESS: double
;             TOPOLOGY: linear
;       MOLECULE TYPE: DNA  (genomic)
;       SEQUENCE DESCRIPTION: SEQ ID NO: 62:
US-09-453-702B-62

  Query Match             57.5%;  Score 575.2;  DB 4;  Length 61663;
  Best Local Similarity   98.2%;  Pred. No. 2.7e-158;
  Matches  603;  Conservative    0;  Mismatches    8;  Indels    3;  Gaps    2;

Qy     389 AAACCCTGTGTGCATCG-TTTTAATTATTCCCGGACACTCCCGCAGAGAAGTTCCCCGTC 447
           |||||| |||||||||| |||| ||||| |||| ||||||||||||||||||||||||||
Db   28589 AAACCCGGTGTGCATCGTTTTTGATTATCCCCGCACACTCCCGCAGAGAAGTTCCCCGTC 28530

Qy     448 AGGGCTGTGGACATAGTTAATCCGGGAATACAATGACGATTCATCGCACCTGACATACAT 507
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28529 AGGGCTGTGGACATAGTTAATCCGGGAATACAATGACGATTCATCGCACCTGACATACAT 28470

Qy     508 TAATAAATATTAACAATATGAAATTTCAACTCATTGTTTAGGGTTTGTTTAATTTTCTAC 567
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28469 TAATAAATATTAACAATATGAAATTTCAACTCATTGTTTAGGGTTTGTTTAATTTTCTAC 28410

Qy     568 ACATACGATTCTGCGAACTTCAAAAAGCATCGGGAATAACACCATGAAAAAAATGCTACT 627
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28409 ACATACGATTCTGCGAACTTCAAAAAGCATCGGGAATAACACCATGAAAAAAATGCTACT 28350

Qy     628 CGCTACTGCGCTGGCCCTGCTTATTACAGGATGTGCTCAACAGACGTTTACTGTTCAAAA 687
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28349 CGCTACTGCGCTGGCCCTGCTTATTACAGGATGTGCTCAACAGACGTTTACTGTTCAAAA 28290

Qy     688 CAAACCGGCAGCAGTAGCACCAAAGGAAACCATCACCCATCATTTCTTCGTTTCTGGAAT 747
           ||||| | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28289 CAAACAGACAGCAGTAGCACCAAAGGAAACCATCACCCATCATTTCTTCGTTTCTGGAAT 28230

Qy     748 TGGGCAGAAGAAAACTGTCGATGCAGCCAAAATTTGTGGCGGCGCAGAAAATGTTGTTAA 807
           |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db   28229 TGGGCAGAAGAAAACTGTCGATGCAGCCAAAATTTGTGGCGGCACAGAAAATGTTGTTAA 28170

Qy     808 AACAGAAACCCAGCAAACATTCGTAAATGGATTGCTCGGTTTTATTACTTTAGGCATTTA 867
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28169 AACAGAAACCCAGCAAACATTCGTAAATGGATTGCTCGGTTTTATTACTTTAGGCATTTA 28110

Qy     868 TACTCCGCTGGAAGCGCGTGTGTATTGCTCACAATAATTGCATGAGTTGCCCATCGCGAT 927
           |||||||||||||||| |||||||||||||||||||||||||||||||  ||||
Db   28109 TACTCCGCTGGAAGCGCGGGTGTATTGCTCACAATAATTGCATGAGTTGCCCAT--CGAT 28052

Qy     928 ATGGGCAACTCTATCTGCACTGCTCATTAATATACTTCTGGGTTCCTTCCAGTTGTTTTT 987
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   28051 ATGGGCAACTCTATCTGCACTGCTCATTAATATACTTCTGGGTTCCTTCCAGTTGTTTTT 27992

Qy     988 GCATAGTGATCAGC 1001
           |||||||||||||
Db   27991 GCATAGTGATCAGC 27978


RESULT 5
US-09-023-221A-6
; Sequence 6, Application US/09023221A
; Patent No. 6087128
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,221A
;      FILING DATE:  12-FEB-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010101
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  6:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  309 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-023-221A-6

  Query Match             30.9%;  Score 309;  DB 3;  Length 309;
  Best Local Similarity   100.0%;  Pred. No. 2.5e-81;
  Matches  309;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy     596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 60

Qy     656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      61 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 120

Qy     716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     121 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 180

Qy     776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     181 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 240

Qy     836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     241 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 300

Qy     896 TCACAATAA 904
           |||||||||
Db     301 TCACAATAA 309
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RESULT 6
US-09-282-352A-6
; Sequence 6, Application US/09282352A
; Patent No. 6187321
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    APPLICANT:  ROBINSON, MICHAEL
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/282,352A
;      FILING DATE:  31-MAR-1999
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 09/023,221
;      FILING DATE:  12-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010102
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  6:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  309 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-282-352A-6

  Query Match             30.9%;  Score 309;  DB 4;  Length 309;
  Best Local Similarity   100.0%;  Pred. No. 2.5e-81;
  Matches  309;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 180

Qy    776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    181 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 240

Qy    836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 300

Qy    896 TCACAATAA 904
          |||||||||
Db    301 TCACAATAA 309
```

```
RESULT 7
US-09-023-221A-1
; Sequence 1, Application US/09023221A
; Patent No. 6087128
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,221A
;      FILING DATE:  12-FEB-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010101
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  760 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-023-221A-1

  Query Match             29.9%;  Score 298.8;  DB 3;  Length 760;
  Best Local Similarity   83.2%;  Pred. No. 3.6e-78;
  Matches  381;  Conservative    0;  Mismatches   62;  Indels   15;  Gaps    3;

Qy    544 TTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGGGAA 603
          |||||   | ||||||||| |   |||||| ||||||||    || || |   ||   | |
Db    243 TTTAGCTATCGTTTAATTATTATCACATAGGATTCTGCCGTTTTTTAACA---ATGCAGGA 299

Qy    604 TAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATGTGC 663
          ||| |  |||||||||||| || |  || | || |||||  ||||||||||||||||||
Db    300 TAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACAGGATGTGC 359

Qy    664 TCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCATCAC 723
          ||||||  ||||||||||  |||||||||| ||||||| ||||||||||||||||||||
Db    360 TCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAAACCATCAC 419

Qy    724 CCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAATTTG 783
          |||||||||||||||| |||||||| ||  |||||||||| ||||||||||||||||||
Db    420 TCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCCAAAATTTG 479

Qy    784 TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATTGCT 843
          ||||||  ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    480 TGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAATGGATTGCT 539

Qy    844 CGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACAATA 903
          |||||||| ||||| ||||| |||||||||||||||| || || ||||||||||||||
Db    540 CGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGCTCACAATA 599

Qy    904 ATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATATACT 963
          |||           |    ||||||||| | ||| |||||||||| ||||||||||||
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Db    600 GTTG----------CCCATCGATATGGGGAGCTC-ATCTGCACTGTTCATTAATATACT 647

Qy    964 TCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          |||||| ||| | ||||||||||||||||||||| |||
Db    648 TCTGGGCTCCCTACAGTTGTTTTTGCATAGTGATAAGC 685


RESULT 8
US-09-282-352A-1
; Sequence 1, Application US/09282352A
; Patent No. 6187321
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    APPLICANT:  ROBINSON, MICHAEL
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/282,352A
;      FILING DATE:  31-MAR-1999
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 09/023,221
;      FILING DATE:  12-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010102
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  760 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-282-352A-1

  Query Match             29.9%;  Score 298.8;  DB 4;  Length 760;
  Best Local Similarity   83.2%;  Pred. No. 3.6e-78;
  Matches  381;  Conservative    0;  Mismatches   62;  Indels   15;  Gaps    3;

Qy    544 TTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGGGAA 603
          |||||   | |||||||| |   |||||| ||||||||     || || |   ||   | |
Db    243 TTTAGCTATCGTTTAATTATTATCACATAGGATTCTGCCGTTTTTAACA---ATGCAGGA 299

Qy    604 TAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATGTGC 663
          ||| |  |||||||||||| || |  || | || |||||  ||||||||||||||||||
Db    300 TAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACAGGATGTGC 359

Qy    664 TCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCATCAC 723
          |||||| ||||||||||||  |||||||||| |||||||| ||||||||||||||||||
Db    360 TCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAAACCATCAC 419

Qy    724 CCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAATTTG 783
           ||||||||||||||||| |||||||| ||  |||||||||| ||||||||||||||||||
Db    420 TCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCCAAAATTTG 479

Qy    784 TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATTGCT 843
          |||||| ||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db    480 TGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAATGGATTGCT 539

Qy    844 CGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACAATA 903
          ||||||||| ||||| ||||| |||||||||||||||||| || || ||||||||||||||
Db    540 CGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGCTCACAATA 599

Qy    904 ATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATATACT 963
          |||            |    ||||||||| | ||| |||||||||| ||||||||||||
Db    600 GTTG----------CCCATCGATATGGGGAGCTC-ATCTGCACTGTTCATTAATATACT 647

Qy    964 TCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          |||||| ||| | ||||||||||||||||||||| |||
Db    648 TCTGGGCTCCCTACAGTTGTTTTTGCATAGTGATAAGC 685


RESULT 9
US-09-023-221A-21
; Sequence 21, Application US/09023221A
; Patent No. 6087128
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,221A
;      FILING DATE:  12-FEB-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010101
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  21:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  378 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-023-221A-21

  Query Match             25.6%;  Score 256.4;  DB 3;  Length 378;
  Best Local Similarity   88.5%;  Pred. No. 7e-66;
  Matches  278;  Conservative    0;  Mismatches   36;  Indels    0;  Gaps    0;

Qy    595 CATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTAC 654
          |||   | |||| |  |||||||||||| || |  || | || ||||| ||||||||||
Db     33 CATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTAC 92

Qy    655 AGGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGA 714
          |||||||||||||| |||||||||||  |||||||||| |||||||| ||||||||||
Db     93 AGGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGA 152

Qy    715 AACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGC 774
          ||||||||| |||||||||||||||| |||||||| ||  |||||||||| ||||||||
Db    153 AACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGC 212
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Qy    775 CAAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          ||||||||||||||| |||||||||||||||||||||||||| ||||||||||||||||||
Db    213 CAAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAA 272

Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
          |||||||||||||||||| ||||| ||||| |||||||||||||||| || || |||||
Db    273 TGGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTG 332

Qy    895 CTCACAATAATTGC 908
          ||||||||| ||||
Db    333 CTCACAATAGTTGC 346


RESULT 10
US-09-282-352A-21
; Sequence 21, Application US/09282352A
; Patent No. 6187321
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    APPLICANT:  ROBINSON, MICHAEL
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/282,352A
;      FILING DATE:  31-MAR-1999
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 09/023,221
;      FILING DATE:  12-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010102
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  21:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  378 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-282-352A-21

  Query Match             25.6%;  Score 256.4;  DB 4;  Length 378;
  Best Local Similarity   88.5%;  Pred. No. 7e-66;
  Matches  278;  Conservative    0;  Mismatches   36;  Indels    0;  Gaps    0;

Qy    595 CATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTAC 654
          |||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db     33 CATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTAC 92

Qy    655 AGGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGA 714
          ||||||||||||||| ||||||||||  |||||||||| |||||||| ||||||||||
Db     93 AGGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGA 152

Qy    715 AACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGC 774
          |||||||| |||||||||||||||| |||||||| ||  |||||||||| ||||||||
Db    153 AACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGC 212

Qy    775 CAAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          ||||||||||||||| ||||||||||||||||||||||||| ||||||||||||||||||
Db    213 CAAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAA 272

Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
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Db    273 TGGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTG 332

Qy    895 CTCACAATAATTGC 908
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Db    333 CTCACAATAGTTGC 346


RESULT 11
US-09-023-221A-22
; Sequence 22, Application US/09023221A
; Patent No. 6087128
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,221A
;      FILING DATE:  12-FEB-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010101
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  22:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  309 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-023-221A-22

  Query Match             25.2%;  Score 252;  DB 3;  Length 309;
  Best Local Similarity   88.6%;  Pred. No. 1.2e-64;
  Matches  273;  Conservative    0;  Mismatches   35;  Indels    0;  Gaps    0;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||||| ||||||||||  |||||||||| |||||||| ||||||||||
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Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
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Db    121 ACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCC 180

Qy    776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
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Db    181 AAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAAT 240

Qy    836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
          |||||||||||||||| ||||| ||||| |||||||||||||||| || || ||||||
Db    241 GGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGC 300

Qy    896 TCACAATA 903
          ||||||||
Db    301 TCACAATA 308


RESULT 12
US-09-282-352A-22
; Sequence 22, Application US/09282352A
; Patent No. 6187321
;  GENERAL INFORMATION:.
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    APPLICANT:  ROBINSON, MICHAEL
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/282,352A
;      FILING DATE:  31-MAR-1999
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 09/023,221
;      FILING DATE:  12-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010102
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  22:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  309 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-282-352A-22

  Query Match             25.2%;  Score 252;  DB 4;  Length 309;
  Best Local Similarity   88.6%;  Pred. No. 1.2e-64;
  Matches  273;  Conservative    0;  Mismatches   35;  Indels    0;  Gaps    0;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||| |||||||||||  |||||||| |||||||| ||||||||||
Db     61 GGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          |||||||| |||||||||||||||| |||||||| ||  |||||||||| |||||||||
Db    121 ACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCC 180

Qy    776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
          |||||||||||||| |||||||||||||||||||||||||| ||||||||||||||||||
Db    181 AAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAAT 240

Qy    836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
          |||||||||||||||| ||||| ||||| |||||||||||||||| || || ||||||
Db    241 GGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGC 300

Qy    896 TCACAATA 903
          ||||||||
Db    301 TCACAATA 308


RESULT 13
US-09-023-221A-5
; Sequence 5, Application US/09023221A
; Patent No. 6087128
;  GENERAL INFORMATION:
;    APPLICANT:  NOLAN, LISA K.
;    APPLICANT:  HORNE, SHELLEY M.
;    TITLE OF INVENTION:  DNA ENCODING AN AVIAN E. COLI ISS
;    NUMBER OF SEQUENCES:  22
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  MUETING, RAASCH & GEBHARDT P.A.
;      STREET:  119 NORTH FOURTH STREET, SUITE 203
;      CITY:  MINNEAPOLIS
;      STATE:  MN
;      COUNTRY:  U.S.A.
;      ZIP:  55401
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,221A
;      FILING DATE:  12-FEB-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  SANDBERG MS., VICTORIA A.
;      REGISTRATION NUMBER:  41,287
;      REFERENCE/DOCKET NUMBER:  255.00010101
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (612) 305-1226
;      TELEFAX:  (612) 305-1228
;  INFORMATION FOR SEQ ID NO:  5:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  309 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
US-09-023-221A-5

  Query Match             22.4%;  Score 224.2;  DB 3;  Length 309;
  Best Local Similarity   87.1%;  Pred. No. 1.7e-56;
  Matches  269;  Conservative    0;  Mismatches   38;  Indels    2;  Gaps    2;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||| |||||||||||  |||||||| |||||||| ||||||||||
Db     61 GGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          |||||||| |||||||||||||||| |||||||| ||  |||||||||| |||||||||
Db    121 ACCATCACTCATCATTTCTTCGTTTCCCCAATTGGACAG-AGAAAACTGTTGATGCAGCC 179

Qy    776 AAAATTTG-TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          |||||||| |||||| |||||||||||||||||||||||||| ||||||||||||||||
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Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
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Db    240 TGCATTGCCCGGTTTTATCACTTTTGGCATCTATACTCCGCGGGAAACCCGTGTATATTG 299

Qy    895 CTCACAATA 903
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Db    300 CTCACAATA 308
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RESULT 14
US-09-282-352A-5
; Sequence 5, Application US/09282352A
; Patent No. 6187321
; GENERAL INFORMATION:
; APPLICANT: NOLAN, LISA K.
; APPLICANT: HORNE, SHELLEY M.
; APPLICANT: ROBINSON, MICHAEL
; TITLE OF INVENTION: DNA ENCODING AN AVIAN E. COLI ISS
; NUMBER OF SEQUENCES: 22
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MUETING, RAASCH & GEBHARDT P.A.
; STREET: 119 NORTH FOURTH STREET, SUITE 203
; CITY: MINNEAPOLIS
; STATE: MN
; COUNTRY: U.S.A.
; ZIP: 55401
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/282,352A
; FILING DATE: 31-MAR-1999
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 09/023,221
; FILING DATE: 12-FEB-1998
; ATTORNEY/AGENT INFORMATION:
; NAME: SANDBERG MS., VICTORIA A.
; REGISTRATION NUMBER: 41,287
; REFERENCE/DOCKET NUMBER: 255.00010102
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (612) 305-1226
; TELEFAX: (612) 305-1228
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 309 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-09-282-352A-5

Query Match 22.4%; Score 224.2; DB 4; Length 309;
Best Local Similarity 87.1%; Pred. No. 1.7e-56;
Matches 269; Conservative 0; Mismatches 38; Indels 2; Gaps 2;

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Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||||| ||||||||||||  |||||||||| |||||||| ||||||||||
Db     61 GGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          |||||||| |||||||||||||||||   |||||| ||| |||||||||| |||||||||
Db    121 ACCATCACTCATCATTTCTTCGTTTCCCCAATTGGACAG-AGAAAACTGTTGATGCAGCC 179

Qy    776 AAAATTTG-TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          |||||||| |||||| |||||||||||||||||||||||||| ||||||||||||||||
Db    180 AAAATTTGTTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAA 239
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Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
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Db    240 TGCATTGCCCGGTTTTATCACTTTTGGCATCTATACTCCGCGGGAAACCCGTGTATATTG 299

Qy    895 CTCACAATA 903
          |||||||||
Db    300 CTCACAATA 308
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RESULT 15
US-09-453-702B-50/c
; Sequence 50, Application US/09453702B
; Patent No. 6365723
; GENERAL INFORMATION:
; APPLICANT: Blattner, Frederick R.
; Burland, Valerie
; Perna, Nicole T.
; Plunkett, Guy
; Welch, Rod
; TITLE OF INVENTION: No. 6365723el Sequences of E. coli O157
; NUMBER OF SEQUENCES: 265
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Quarles & Brady
; STREET: 1 South Pinckney Street
; CITY: Madison
; STATE: WI
; COUNTRY: US
; ZIP: 53701-2113
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Word Perfect 8.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/453,702B
; FILING DATE: 03-Dec-1999
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/110,955
; FILING DATE: 04-DEC-1998
; ATTORNEY/AGENT INFORMATION:
; NAME: Seay, Nicholas J.
; REGISTRATION NUMBER: 27386
; REFERENCE/DOCKET NUMBER: 960296.95017
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (608) 251-5000
; TELEFAX: (608) 251-9166
; INFORMATION FOR SEQ ID NO: 50:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 38584
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
; SEQUENCE DESCRIPTION: SEQ ID NO: 50:
US-09-453-702B-50

Query Match 5.9%; Score 59.2; DB 4; Length 38584;
Best Local Similarity 77.8%; Pred. No. 3.6e-07;
Matches 84; Conservative 0; Mismatches 23; Indels 1; Gaps 1;

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Qy    448 AGGGCTGTGGACATAGTTAATCCGGGAATACAATGACGATTCATCGCA 495
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(without alignments)
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Title: US-09-740-211-13_COPY_8700_9700
Perfect score: 1001
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 381593 seqs, 216252194 residues

Total number of hits satisfying chosen parameters: 763186

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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14: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
	1	1001	100.0	11933	9	US-10-007-968-13	Sequence 13, Appl
	2	1001	100.0	11933	10	US-09-740-211-13	Sequence 13, Appl
	3	309	30.9	309	10	US-09-738-599-6	Sequence 6, Appli
	4	298.8	29.9	760	10	US-09-738-599-1	Sequence 1, Appli
	5	256.4	25.6	378	10	US-09-738-599-21	Sequence 21, Appl
	6	252	25.2	309	10	US-09-738-599-22	Sequence 22, Appl
	7	224.2	22.4	309	10	US-09-738-599-5	Sequence 5, Appli
	8	40.8	4.1	627	10	US-09-728-446-280	Sequence 280, App
	9	40.8	4.1	3758	9	US-10-108-605-118	Sequence 118, App
	10	38.4	3.8	361	10	US-09-960-352-7535	Sequence 7535, Ap
c	11	35.8	3.6	302	10	US-09-969-373-413	Sequence 413, App
c	12	35.8	3.6	302	10	US-09-969-373-417	Sequence 417, App
	13	35.8	3.6	1823	9	US-09-938-842A-4744	Sequence 4744, Ap
c	14	35.8	3.6	2000	9	US-09-938-842A-3589	Sequence 3589, Ap
	15	35.6	3.6	25950	10	US-09-764-870-597	Sequence 597, App
	16	35	3.5	1038	9	US-09-938-842A-3994	Sequence 3994, Ap
	17	35	3.5	1040	10	US-09-887-576-696	Sequence 696, App
c	18	35	3.5	6945	10	US-09-764-887-314	Sequence 314, App
	19	35	3.5	7657	10	US-09-070-927A-33	Sequence 33, Appl
	20	35	3.5	8354	9	US-10-125-818-1	Sequence 1, Appli
c	21	34.8	3.5	380	10	US-09-864-761-3795	Sequence 3795, Ap
	22	34.6	3.5	578	10	US-09-815-242-2027	Sequence 2027, Ap
c	23	34.6	3.5	669	10	US-09-815-242-8992	Sequence 8992, Ap
c	24	34.6	3.5	672	10	US-09-815-242-4834	Sequence 4834, Ap
c	25	34.6	3.5	2732	10	US-09-771-161A-90	Sequence 90, Appl
c	26	34.4	3.4	492	9	US-09-796-692-7148	Sequence 7148, Ap
	27	34.4	3.4	633	10	US-09-861-451A-37	Sequence 37, Appl
	28	34.2	3.4	432	10	US-09-960-352-14612	Sequence 14612, A
c	29	34.2	3.4	479	10	US-09-783-590-4423	Sequence 4423, Ap
c	30	34	3.4	1181	9	US-09-938-842A-4549	Sequence 4549, Ap
c	31	34	3.4	684973	10	US-09-263-959-1	Sequence 1, Appli
c	32	33.8	3.4	21784	10	US-09-820-002-3	Sequence 3, Appli
	33	33.8	3.4	172637	10	US-09-805-458A-3	Sequence 3, Appli
c	34	33.6	3.4	444	10	US-09-960-352-13335	Sequence 13335, A
	35	33.6	3.4	1817	10	US-09-452-599-1	Sequence 1, Appli
c	36	33.6	3.4	2463	10	US-09-842-552-106	Sequence 106, App
c	37	33.6	3.4	3280	10	US-09-842-552-80	Sequence 80, Appl
c	38	33.6	3.4	4951	10	US-09-070-927A-261	Sequence 261, App
	39	33.4	3.3	483	10	US-09-880-107-2669	Sequence 2669, Ap
c	40	33.4	3.3	1325	10	US-09-765-205-1	Sequence 1, Appli
c	41	33.4	3.3	1333	9	US-09-978-295A-321	Sequence 321, App
c	42	33.4	3.3	1333	9	US-09-978-697-321	Sequence 321, App
c	43	33.4	3.3	1333	9	US-09-978-192A-321	Sequence 321, App
c	44	33.4	3.3	1333	9	US-09-999-832A-321	Sequence 321, App
c	45	33.4	3.3	1391	12	US-10-044-477-2	Sequence 2, Appli

ALIGNMENTS

RESULT 1
US-10-007-968-13
; Sequence 13, Application US/10007968
; Patent No. US20020159977A1
; GENERAL INFORMATION:
; APPLICANT: Couto, Linda B.
; APPLICANT: Colosi, Peter C.
; TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
; TITLE OF INVENTION: by Target Cells
; FILE REFERENCE: Avigen-04082
; CURRENT APPLICATION NUMBER: US/10/007,968
; CURRENT FILING DATE: 2001-12-13
; PRIOR APPLICATION NUMBER: 09/740,211
; PRIOR FILING DATE: 2000-12-18
; PRIOR APPLICATION NUMBER: 60/125,974
; PRIOR FILING DATE: 1999-03-24
; PRIOR APPLICATION NUMBER: 60/104,994
; PRIOR FILING DATE: 1998-10-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
; LENGTH: 11933
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-007-968-13

Query Match 100.0%; Score 1001; DB 9; Length 11933;
Best Local Similarity 100.0%; Pred. No. 4.6e-251;
Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy       1 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8700 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 8759

Qy      61 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8760 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 8819

Qy     121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    8820 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 8879

Qy     181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8880 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 8939

Qy     241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8940 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 8999

Qy     301 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGGCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9000 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 9059

Qy     361 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9060 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 9119

Qy     421 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9120 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 9179

Qy     481 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9180 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 9239

Qy     541 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9240 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 9299

Qy     601 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9300 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 9359

Qy     661 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9360 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 9419

Qy     721 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9420 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 9479

Qy     781 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9480 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 9539

Qy     841 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 900
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9540 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 9599

Qy     901 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 960
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9600 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 9659

Qy     961 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
           |||||||||||||||||||||||||||||||||||||||||
Db    9660 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 9700


RESULT 2
US-09-740-211-13
; Sequence 13, Application US/09740211
; Patent No. US20010010815A1
; GENERAL INFORMATION:
;  APPLICANT: Couto, Linda B.
;  APPLICANT:  Colosi, Peter C.
;  TITLE OF INVENTION: Adeno-Associated Vectors for Expression of Factor VIII
;  TITLE OF INVENTION:  by Target Cells
;  FILE REFERENCE: Avigen-04082
;  CURRENT APPLICATION NUMBER: US/09/740,211
;  CURRENT FILING DATE:  2000-12-18
;  PRIOR APPLICATION NUMBER: 09/470,618
;  PRIOR FILING DATE: 1999-12-22
;  PRIOR APPLICATION NUMBER: 60/125,974
;  PRIOR FILING DATE: 1999-03-24
;  PRIOR APPLICATION NUMBER: 60/104,994
;  PRIOR FILING DATE: 1998-10-20
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
;   LENGTH: 11933
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence:  Synthetic
US-09-740-211-13

  Query Match             100.0%;  Score 1001;  DB 10;  Length 11933;
  Best Local Similarity   100.0%;  Pred. No. 4.6e-251;
  Matches 1001;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy       1 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8700 AACGTCTCACATTATATTTACTATCTAGCCACAGATAATATTCACATCGTGTTAGAAAAC 8759

Qy      61 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8760 GATAACACCGTGTTAATAAAAGGACTTAAAAAGGTTGTAAATGTTAAATTCTCAAGAAAC 8819

Qy     121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8820 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 8879

Qy     181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8880 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 8939

Qy     241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    8940 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 8999

Qy     301 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGGCG 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9000 ACACAAATCTTTCCACGCTAAATCATAACGTCCGGTTTCTTCCGTGTCAGCACCGGGGCG 9059

Qy     361 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9060 TTGGCATAATGCAATACGTGTACGCGCTAAACCCTGTGTGCATCGTTTTAATTATTCCCG 9119

Qy     421 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9120 GACACTCCCGCAGAGAAGTTCCCCGTCAGGGCTGTGGACATAGTTAATCCGGGAATACAA 9179

Qy     481 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9180 TGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGAAATTTCAACTCA 9239

Qy     541 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9240 TTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGG 9299

Qy     601 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 660
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9300 GAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATG 9359

Qy     661 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9360 TGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCAT 9419

Qy     721 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 780
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9420 CACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAAT 9479

Qy     781 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 840
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    9480 TTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATT 9539
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Qy    841 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9540 GCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACA 9599

Qy    901 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9600 ATAATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATAT 9659

Qy    961 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          |||||||||||||||||||||||||||||||||||||||
Db   9660 ACTTCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 9700


RESULT 3
US-09-738-599-6
; Sequence 6, Application US/09738599
; Patent No. US20020102546A1
; GENERAL INFORMATION:
;  APPLICANT: No. US20020102546Alan, Lisa
;  APPLICANT:  Horne, Shelley
;  TITLE OF INVENTION: NUCLEIC ACID ENCODING AN AVIAN E. COLI ISS POLYPEPTIDE AND METHOD
;  FILE REFERENCE: 255.0001 0122
;  CURRENT APPLICATION NUMBER: US/09/738,599
;  CURRENT FILING DATE:  2000-12-15
;  PRIOR APPLICATION NUMBER: 09/282,352
;  PRIOR FILING DATE: 1999-03-31
;  PRIOR APPLICATION NUMBER: 09/023,221
;  PRIOR FILING DATE: 1998-02-12
;  NUMBER OF SEQ ID NOS: 26
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 6
;   LENGTH: 309
;   TYPE: DNA
;   ORGANISM: Bacteriophage lambda
US-09-738-599-6

  Query Match             30.9%;  Score 309;  DB 10;  Length 309;
  Best Local Similarity   100.0%;  Pred. No. 2.2e-71;
  Matches  309;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 180

Qy    776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    181 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 240

Qy    836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 300

Qy    896 TCACAATAA 904
          |||||||||
Db    301 TCACAATAA 309


RESULT 4
US-09-738-599-1
; Sequence 1, Application US/09738599
; Patent No. US20020102546A1
; GENERAL INFORMATION:
;  APPLICANT: No. US20020102546Alan, Lisa
;  APPLICANT:  Horne, Shelley
;  TITLE OF INVENTION: NUCLEIC ACID ENCODING AN AVIAN E. COLI ISS POLYPEPTIDE AND MET
;  FILE REFERENCE: 255.0001 0122
;  CURRENT APPLICATION NUMBER: US/09/738,599
;  CURRENT FILING DATE:  2000-12-15
;  PRIOR APPLICATION NUMBER: 09/282,352
;  PRIOR FILING DATE: 1999-03-31
;  PRIOR APPLICATION NUMBER: 09/023,221
;  PRIOR FILING DATE: 1998-02-12
;  NUMBER OF SEQ ID NOS: 26
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 1
;   LENGTH: 760
;   TYPE: DNA
;   ORGANISM: Escherichia coli
US-09-738-599-1

  Query Match             29.9%;  Score 298.8;  DB 10;  Length 760;
  Best Local Similarity   83.2%;  Pred. No. 1.4e-68;
  Matches  381;  Conservative    0;  Mismatches   62;  Indels   15;  Gaps    3;

Qy    544 TTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGGGAA 603
          |||||   | |||||||| |   |||||| ||||||||    || || |   ||   | |
Db    243 TTTAGCTATCGTTTAATTATTATCACATAGGATTCTGCCGTTTTTAACA---ATGCAGGA 299

Qy    604 TAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACAGGATGTGC 663
          ||| |  |||||||||||| || |  || | || |||||  ||||||||||||||||||
Db    300 TAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACAGGATGTGC 359

Qy    664 TCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAAACCATCAC 723
          |||||| ||||||||||||  |||||||| |||||||| ||||||||||||||||||||
Db    360 TCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAAACCATCAC 419

Qy    724 CCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCCAAAATTTG 783
           ||||||||||||||||| |||||||| ||  |||||||||| ||||||||||||||||||
Db    420 TCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCCAAAATTTG 479

Qy    784 TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAATGGATTGCT 843
          |||||| ||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db    480 TGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAATGGATTGCT 539

Qy    844 CGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGCTCACAATA 903
          ||||||||| ||||| ||||| |||||||||||||||||| || || ||||||||||||||
Db    540 CGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGCTCACAATA 599

Qy    904 ATTGCATGAGTTGCCCATCGCGATATGGGCAACTCTATCTGCACTGCTCATTAATATACT 963
          |||          |    |||||||| | ||| ||||||||| ||||||||||||||
Db    600 GTTG-----------CCCATCGATATGGGGAGCTC-ATCTGCACTGTTCATTAATATACT 647

Qy    964 TCTGGGTTCCTTCCAGTTGTTTTTGCATAGTGATCAGC 1001
          |||||| ||| | |||||||||||||||||||| |||
Db    648 TCTGGGCTCCCTACAGTTGTTTTTGCATAGTGATAAGC 685


RESULT 5
US-09-738-599-21
; Sequence 21, Application US/09738599
; Patent No. US20020102546A1
; GENERAL INFORMATION:
;  APPLICANT: No. US20020102546Alan, Lisa
;  APPLICANT:  Horne, Shelley
;  TITLE OF INVENTION: NUCLEIC ACID ENCODING AN AVIAN E. COLI ISS POLYPEPTIDE AND MET
;  FILE REFERENCE: 255.0001 0122
;  CURRENT APPLICATION NUMBER: US/09/738,599
;  CURRENT FILING DATE:  2000-12-15
;  PRIOR APPLICATION NUMBER: 09/282,352
;  PRIOR FILING DATE: 1999-03-31
;  PRIOR APPLICATION NUMBER: 09/023,221
;  PRIOR FILING DATE: 1998-02-12
;  NUMBER OF SEQ ID NOS: 26
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 21
;   LENGTH: 378
;   TYPE: DNA
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;   ORGANISM: Escherichia coli
US-09-738-599-21

  Query Match             25.6%;  Score 256.4;  DB 10;  Length 378;
  Best Local Similarity   88.5%;  Pred. No. 1.2e-57;
  Matches  278;  Conservative    0;  Mismatches   36;  Indels    0;  Gaps    0;

Qy    595 CATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTAC 654
          |||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db     33 CATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTAC 92

Qy    655 AGGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGA 714
          ||||||||||||| ||||||||||  |||||||||| |||||||| ||||||||||
Db     93 AGGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGA 152

Qy    715 AACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGC 774
          |||||||| |||||||||||||||| |||||||| ||  |||||||||| ||||||||
Db    153 AACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGC 212

Qy    775 CAAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          |||||||||||||| |||||||||||||||||||||||||||| ||||||||||||||||
Db    213 CAAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAA 272

Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
          |||||||||||||||| ||||| ||||| |||||||||||||||| || || |||||
Db    273 TGGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTG 332

Qy    895 CTCACAATAATTGC 908
          ||||||||| ||||
Db    333 CTCACAATAGTTGC 346


RESULT 6
US-09-738-599-22
; Sequence 22, Application US/09738599
; Patent No. US20020102546A1
; GENERAL INFORMATION:
;  APPLICANT: No. US20020102546A1an, Lisa
;  APPLICANT:  Horne, Shelley
;  TITLE OF INVENTION: NUCLEIC ACID ENCODING AN AVIAN E. COLI ISS POLYPEPTIDE AND METHOD
;  FILE REFERENCE: 255.0001 0122
;  CURRENT APPLICATION NUMBER: US/09/738,599
;  CURRENT FILING DATE:  2000-12-15
;  PRIOR APPLICATION NUMBER: 09/282,352
;  PRIOR FILING DATE: 1999-03-31
;  PRIOR APPLICATION NUMBER: 09/023,221
;  PRIOR FILING DATE: 1998-02-12
;  NUMBER OF SEQ ID NOS: 26
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 22
;   LENGTH: 309
;   TYPE: DNA
;   ORGANISM: Escherichia coli
US-09-738-599-22

  Query Match             25.2%;  Score 252;  DB 10;  Length 309;
  Best Local Similarity   88.6%;  Pred. No. 1.6e-56;
  Matches  273;  Conservative    0;  Mismatches   35;  Indels    0;  Gaps    0;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          |||||||||||||| ||||||||||  |||||||||| |||||||| ||||||||||
Db     61 GGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          |||||||| |||||||||||||||| |||||||| ||  |||||||||| |||||||||
Db    121 ACCATCACTCATCATTTCTTCGTTTCGGGAATTGGACAAGAGAAAACTGTTGATGCAGCC 180

Qy    776 AAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAAT 835
          |||||||||||||| |||||||||||||||||||||||||||| ||||||||||||||||
Db    181 AAAATTTGTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAAT 240

Qy    836 GGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTGC 895
          |||||||||||||||| ||||| ||||| |||||||||||||||| || || ||||||
Db    241 GGATTGCTCGGTTTTATCACTTTTGGCATCTATACTCCGCTGGAAGCCCGGGTATATTGC 300

Qy    896 TCACAATA 903
          ||||||||
Db    301 TCACAATA 308


RESULT 7
US-09-738-599-5
; Sequence 5, Application US/09738599
; Patent No. US20020102546A1
; GENERAL INFORMATION:
;  APPLICANT: No. US20020102546A1an, Lisa
;  APPLICANT:  Horne, Shelley
;  TITLE OF INVENTION: NUCLEIC ACID ENCODING AN AVIAN E. COLI ISS POLYPEPTIDE AND MET
;  FILE REFERENCE: 255.0001 0122
;  CURRENT APPLICATION NUMBER: US/09/738,599
;  CURRENT FILING DATE:  2000-12-15
;  PRIOR APPLICATION NUMBER: 09/282,352
;  PRIOR FILING DATE: 1999-03-31
;  PRIOR APPLICATION NUMBER: 09/023,221
;  PRIOR FILING DATE: 1998-02-12
;  NUMBER OF SEQ ID NOS: 26
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 5
;   LENGTH: 309
;   TYPE: DNA
;   ORGANISM: Escherichia coli
US-09-738-599-5

  Query Match             22.4%;  Score 224.2;  DB 10;  Length 309;
  Best Local Similarity   87.1%;  Pred. No. 2.7e-49;
  Matches  269;  Conservative    0;  Mismatches   38;  Indels    2;  Gaps    2;

Qy    596 ATCGGGAATAACACCATGAAAAAAATGCTACTCGCTACTGCGCTGGCCCTGCTTATTACA 655
          ||   | |||| |  |||||||||||| || |  || | || |||||  ||||||||||
Db      1 ATGCAGGATAATAAGATGAAAAAAATGTTATTTTCTGCCGCTCTGGCAATGCTTATTACA 60

Qy    656 GGATGTGCTCAACAGACGTTTACTGTTCAAAACAAACCGGCAGCAGTAGCACCAAAGGAA 715
          |||||||||||||| ||||||||||  |||||||||| |||||||| ||||||||||
Db     61 GGATGTGCTCAACAAACGTTTACTGTTGGAAACAAACCGACAGCAGTAACACCAAAGGAA 120

Qy    716 ACCATCACCCATCATTTCTTCGTTTCTGGAATTGGGCAGAAGAAAACTGTCGATGCAGCC 775
          |||||||| |||||||||||||||||   |||||| ||| |||||||||| |||||||||
Db    121 ACCATCACTCATCATTTCTTCGTTTCCCCAATTGGACAG-AGAAAACTGTTGATGCAGCC 179

Qy    776 AAAATTTG-TGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCCAGCAAACATTCGTAAA 834
          |||||||| |||||| |||||||||||||||||||||||||| ||||||||||||||||
Db    180 AAAATTTGTTGGCGGTGCAGAAAATGTTGTTAAAACAGAAACTCAGCAAACATTCGTAAA 239

Qy    835 TGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGGAAGCGCGTGTGTATTG 894
          || ||||| |||||||||| ||||| ||||| |||||||||| |||| | ||||| |||||
Db    240 TGCATTGCCCGGTTTTATCACTTTTGGCATCTATACTCCGCGGGAAACCCGTGTATATTG 299

Qy    895 CTCACAATA 903
          |||||||||
Db    300 CTCACAATA 308


RESULT 8
US-09-728-446-280
; Sequence 280, Application US/09728446
; Patent No. US20020081668A1
; GENERAL INFORMATION:
;  APPLICANT: Friedrich, Glenn
;  APPLICANT:  Zambrowicz, Brian
;  APPLICANT:  Sands, Arthur T.
;  TITLE OF INVENTION: No. US20020081668A1el Murine Polynucleotide Sequences
;  TITLE OF INVENTION:  and Mutant Cells and Mutant Animals Defined Thereby
```

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;  FILE REFERENCE: LEX-0101-USA
;  CURRENT APPLICATION NUMBER: US/09/728,446
;  CURRENT FILING DATE:  2000-11-30
;  PRIOR APPLICATION NUMBER: US 60/168,270
;  PRIOR FILING DATE: 1999-12-01
;  NUMBER OF SEQ ID NOS: 1461
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 280
;   LENGTH: 627
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(627)
;   OTHER INFORMATION: n = A,T,C or G
US-09-728-446-280

  Query Match             4.1%;  Score 40.8;  DB 10;  Length 627;
  Best Local Similarity   52.3%;  Pred. No. 0.22;
  Matches   90;  Conservative    0;  Mismatches   82;  Indels    0;  Gaps    0;

Qy    759 AAACTGTCGATGCAGCCAAAATTTGTGGCGGCGCAGAAAATGTTGTTAAAACAGAAACCC 818
          ||  | | | |  |||  | || ||    |   ||| |       ||| ||     ||||
Db    256 AATATTTTGCTAAAGCACACATGTGAAAGGATACAGGATGAAGGCTTATAAATATGACCC 315

Qy    819 AGCAAACATTCGTAAATGGATTGCTCGGTTTTATTACTTTAGGCATTTATACTCCGCTGG 878
          | |||||| | | |  | ||  |||   ||   || ||  |    | ||| || |||||
Db    316 AACAAACAATGGGAGCTCGAGCGCTGCTTTGCTTTGCTCCACCTCTCTATTCTTCGCTGA 375

Qy    879 AAGCGCGTGTGTATTGCTCACAATAATTGCATGAGTTGCCCATCGCGATATG 930
           | | | |||||    || ||  |   ||  |||||| | | |   |  |||
Db    376 CAACACATGTGTTCGTCTTACGTTGCGTGGCTGAGTTCCACTTGTGGTGATG 427


RESULT 9
US-10-108-605-118
; Sequence 118, Application US/10108605
; Patent No. US20020160934A1
; GENERAL INFORMATION:
;  APPLICANT: Broadus, Julie
;  APPLICANT:  Stam, Lynn
;  APPLICANT:  Bachmann, Jane
;  APPLICANT:  Kamdar, Kim
;  TITLE OF INVENTION: NUCLEIC ACID SEQUENCES FROM DROSOPHILA MELANOGASTER THAT ENCODE
;  TITLE OF INVENTION:  PROTEINS ESSENTIAL FOR LARVAL VIABILITY AND USES THEREOF
;  FILE REFERENCE: 31133B
;  CURRENT APPLICATION NUMBER: US/10/108,605
;  CURRENT FILING DATE:  2002-03-27
;  PRIOR APPLICATION NUMBER: US 09/761,142
;  PRIOR FILING DATE: 2001-01-16
;  PRIOR APPLICATION NUMBER: US 60/176,418
;  PRIOR FILING DATE: 2000-01-14
;  NUMBER OF SEQ ID NOS: 361
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 118
;   LENGTH: 3758
;   TYPE: DNA
;   ORGANISM: Drosophila melanogaster
US-10-108-605-118

  Query Match             4.1%;  Score 40.8;  DB 9;  Length 3758;
  Best Local Similarity   50.0%;  Pred. No. 0.44;
  Matches  102;  Conservative    0;  Mismatches  102;  Indels    0;  Gaps    0;

Qy    121 ACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAA 180
          | || ||||  |    |     ||| |||  |||  |  ||   ||| | |||||   ||
Db   2657 AGGCCTCTTTCAATTTCTAGGGATGGTAGCGTGATCTTTAGTATAATAAAATTTCTTAAA 2716

Qy    181 TACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAACAT 240
          |||     |||   || ||  |        ||  | ||  ||  ||  |||    | |  ||
Db   2717 TACTTATCAAAAAGTGATACGGATAAGTATTCTCTATCTGTATCAGTTATTTTTAATTAT 2776

Qy    241 AAAAGATATTACTATACCTTTGATAATTCATTACTATTTACTGAGAGCATTCAGAACACT 300
           ||| |||    ||| | ||| |||   | | || |   | | |  | ||   | |||
Db   2777 TTCAGAAATTGTCCTACTTCTGAAAATATTTAAGTAGTATATAAAAATAATCTATATACT 2836

Qy    301 ACACAAATCTTTCCACGCTAAATC 324
            ||| | |||  |    ||| |
Db   2837 TTGCAATTTTTTTTATTTAAAAAC 2860


RESULT 10
US-09-960-352-7535
; Sequence 7535, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
;  APPLICANT: Warren, Wesley C.
;  APPLICANT:  Tao, Nengbing
;  APPLICANT:  Byatt, John C.
;  APPLICANT:  Mathialagan, Nagappan
;  TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
;  TITLE OF INVENTION:  MUSCLE AND FAT DEPOSITION
;  FILE REFERENCE: 16511.006/37-21(10298)C
;  CURRENT APPLICATION NUMBER: US/09/960,352
;  CURRENT FILING DATE:  2001-09-24
;  NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 7535
;   LENGTH: 361
;   TYPE: DNA
;   ORGANISM: Bos taurus
;   OTHER INFORMATION: Clone ID: 32-LIB34-056-Q1-E1-H7
US-09-960-352-7535

  Query Match             3.8%;  Score 38.4;  DB 10;  Length 361;
  Best Local Similarity   51.8%;  Pred. No. 0.75;
  Matches   87;  Conservative    0;  Mismatches   81;  Indels    0;  Gaps    0;

Qy    143 ATGATAGGTTGAAATCAAGAGAAATCACATTTCAGCAATACAGGGAAAATCTTGCTAAAG 202
          ||||  |    ||  | |   ||||      || || | | ||| | |||   | ||||
Db     86 ATGACCGTGAAAATGCCACCAAAATTCTCCATCGGCCAAAGAGGTACAATTCAGGTAAAT 145

Qy    203 CAGGAGTTTTCCGATGGGTTACAAATATCCATGAACATAAAAGATATTACTATACCTTTG 262
            | ||  ||     | |  |    |      |||  |||| | |  | | ||| ||||
Db    146 TGGAAGAGTTTGTTCGAGGGAACCTTGAGAGAGAATGTAAAGAAGAAAAATGTACTTTTG 205

Qy    263 ATAATTCATTACTATTTACTGAGAGCATTCAGAACACTACACAAATCT 310
          |  |  ||  || | ||  ||| | || | |||| |||||  || | |
Db    206 AAGAAGCACGACAAGTTTTTGAAAACACTGAGAAAACTACTGAATTTT 253


RESULT 11
US-09-969-373-413/c
; Sequence 413, Application US/09969373
; Patent No. US20020133852A1
; GENERAL INFORMATION:
;  APPLICANT: Effertz, Roger J.
;  APPLICANT:  Hauge, Brian M.
;  TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
;  FILE REFERENCE: 38-10(52679)A
;  CURRENT APPLICATION NUMBER: US/09/969,373
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 09/754,853
;  PRIOR FILING DATE: 2001-01-05
;  PRIOR APPLICATION NUMBER: US 09/760,427
;  PRIOR FILING DATE: 2001-01-13
;  PRIOR APPLICATION NUMBER: US 09/855,768
;  PRIOR FILING DATE: 2001-05-15
;  NUMBER OF SEQ ID NOS: 4593.
; SEQ ID NO 413
;   LENGTH: 302
;   TYPE: DNA
;   ORGANISM: Glycine max
US-09-969-373-413

  Query Match             3.6%;  Score 35.8;  DB 10;  Length 302;
  Best Local Similarity   55.1%;  Pred. No. 3.3;
```

```
  Matches   70;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

Qy     501 CATACATTAATAAATATTAACAATATGAAATTTCAACTCATTGTTTAGGGTTTGTTTAAT 560
           |||    ||| | | |       | |   | | | |  | ||| ||||  ||    |||||
Db     272 CATCATGTAAGAGAAAAATCATAAAATTATTATTATTTAATTTTTTAATGTAAAGTTAAT 213

Qy     561 TTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGGGAATAACACCATGAAAAAAA 620
           ||| | ||| ||   | ||   ||  ||||||   ||| | || || | |||  | |||
Db     212 TTTTTCCACTTATATTCCTTATAATATCAAAACCGATCAGTAACAAAATCATTCATAAAT 153

Qy     621 TGCTACT 627
           |  || |
Db     152 TAATAAT 146


RESULT 12
US-09-969-373-417/c
; Sequence 417, Application US/09969373
; Patent No. US20020133852A1
; GENERAL INFORMATION:
;  APPLICANT: Effertz, Roger J.
;  APPLICANT:  Hauge, Brian M.
;  TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
;  FILE REFERENCE: 38-10(52679)A
;  CURRENT APPLICATION NUMBER: US/09/969,373
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 09/754,853
;  PRIOR FILING DATE: 2001-01-05
;  PRIOR APPLICATION NUMBER: US 09/760,427
;  PRIOR FILING DATE: 2001-01-13
;  PRIOR APPLICATION NUMBER: US 09/855,768
;  PRIOR FILING DATE: 2001-05-15
;  NUMBER OF SEQ ID NOS: 4593
; SEQ ID NO 417
;   LENGTH: 302
;   TYPE: DNA
;   ORGANISM: Glycine max
US-09-969-373-417

  Query Match             3.6%;  Score 35.8;  DB 10;  Length 302;
  Best Local Similarity   55.1%;  Pred. No. 3.3;
  Matches   70;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

Qy     501 CATACATTAATAAATATTAACAATATGAAATTTCAACTCATTGTTTAGGGTTTGTTTAAT 560
           |||    ||| | | |       | |   | | | |  | ||| ||||  ||    |||||
Db     272 CATCATGTAAGAGAAAAATCATAAAATTATTATTATTTAATTTTTTAATGTAAAGTTAAT 213

Qy     561 TTTCTACACATACGATTCTGCGAACTTCAAAAAGCATCGGGAATAACACCATGAAAAAAA 620
           ||| | ||| ||   | ||   ||  ||||||   ||| | || || | |||  | |||
Db     212 TTTTTCCACTTATATTCCTTATAATATCAAAACCGATCAGTAACAAAATCATTCATAAAT 153

Qy     621 TGCTACT 627
           |  || |
Db     152 TAATAAT 146


RESULT 13
US-09-938-842A-4744
; Sequence 4744, Application US/09938842A
; Patent No. US20020160378A1
; GENERAL INFORMATION:
;  APPLICANT: Harper, Jeff
;  APPLICANT:  Kreps, Joel
;  APPLICANT:  Wang, Xun
;  APPLICANT:  Zhu, Tong
;  TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
;  TITLE OF INVENTION:  SAME, AND METHODS OF USE
;  FILE REFERENCE: SCRIP1300-3
;  CURRENT APPLICATION NUMBER: US/09/938,842A
;  CURRENT FILING DATE:  2001-08-24
;  PRIOR APPLICATION NUMBER: US 60/227,866
;  PRIOR FILING DATE: 2000-08-24
;  PRIOR APPLICATION NUMBER: US 60/264,647
;  PRIOR FILING DATE: 2001-01-16
;  PRIOR APPLICATION NUMBER: US 60/300,111
;  PRIOR FILING DATE: 2001-06-22
;  NUMBER OF SEQ ID NOS: 5379
; SEQ ID NO 4744
;   LENGTH: 1823
;   TYPE: DNA
;   ORGANISM: Arabidopsis thaliana
US-09-938-842A-4744

  Query Match             3.6%;  Score 35.8;  DB 9;  Length 1823;
  Best Local Similarity   53.1%;  Pred. No. 6.7;
  Matches   76;  Conservative    0;  Mismatches   67;  Indels    0;  Gaps    0;

Qy     216 ATGGGTTACAAATATCCATGAACATAAAAGATATTACTATACCTTTGATAATTCATTACT 275
           |||  || || || | |||| | ||  || || |   |||   ||| | ||||||   ||
Db     478 ATGTTTTTCACATCTTCATGTATATTTAACATTTATTTATTTTTTAAAAATTCAACTCT 537

Qy     276 ATTTACTGAGAGCATTCAGAACACTACACAAATCTTTCCACGCTAAATCATAACGTCCGG 335
           |||    || |   |   || | || |  ||| |     ||  ||||| | ||
Db     538 TTTTTTAAAGGGTTATATAAATATTAAATTAATATGATGTCGTGAAATCTAACCGGATTT 597

Qy     336 TTTCTTCCGTGTCAGCACCGGGG 358
           |   | |||     | |||| ||
Db     598 TGGATCCCGCCATTGAACCGTGG 620


RESULT 14
US-09-938-842A-3589/c
; Sequence 3589, Application US/09938842A
; Patent No. US20020160378A1
; GENERAL INFORMATION:
;  APPLICANT: Harper, Jeff
;  APPLICANT:  Kreps, Joel
;  APPLICANT:  Wang, Xun
;  APPLICANT:  Zhu, Tong
;  TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
;  TITLE OF INVENTION:  SAME, AND METHODS OF USE
;  FILE REFERENCE: SCRIP1300-3
;  CURRENT APPLICATION NUMBER: US/09/938,842A
;  CURRENT FILING DATE:  2001-08-24
;  PRIOR APPLICATION NUMBER: US 60/227,866
;  PRIOR FILING DATE: 2000-08-24
;  PRIOR APPLICATION NUMBER: US 60/264,647
;  PRIOR FILING DATE: 2001-01-16
;  PRIOR APPLICATION NUMBER: US 60/300,111
;  PRIOR FILING DATE: 2001-06-22
;  NUMBER OF SEQ ID NOS: 5379
; SEQ ID NO 3589
;   LENGTH: 2000
;   TYPE: DNA
;   ORGANISM: Arabidopsis thaliana
US-09-938-842A-3589

  Query Match             3.6%;  Score 35.8;  DB 9;  Length 2000;
  Best Local Similarity   53.1%;  Pred. No. 7;
  Matches   76;  Conservative    0;  Mismatches   67;  Indels    0;  Gaps    0;

Qy     469 CCGGGAATACAATGACGATTCATCGCACCTGACATACATTAATAAATATTAACAATATGA 528
           ||| | | | |   |  ||| ||      | | | |   | ||||||  |||||  | ||
Db     267 CCGAGGAAAAAGCAAATATTTATTATTTATAAGAAAATATTATAAATTAAAACAAAAAGA 208

Qy     529 AATTTCAACTCATTGTTTAGGGTTTGTTTAATTTTCTACACATACGATTCTGCGAACTTC 588
           ||    ||   |||| || |    ||| || | |  | | |   |||| |   ||  ||
Db     207 AAAAAAAAAAAATTGAATATGAAGAGTTAAAATATGAAAAAAGTGGATTTTCGTAAGATC 148

Qy     589 AAAAAGCATCGGGAATAACACCA 611
           ||| ||| ||   |  || || |
Db     147 AAATAGCGTCAAAATCAAGACAA 125


RESULT 15
US-09-764-870-597
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; Sequence 597, Application US/09764870
; Patent No. US20020042386A1
; GENERAL INFORMATION:
;  APPLICANT: Rosen et al.
;  TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
;  FILE REFERENCE: PTZ14
;  CURRENT APPLICATION NUMBER: US/09/764,870
;  CURRENT FILING DATE:  2001-01-17
;  Prior application data removed - consult PALM or file wrapper
;  NUMBER OF SEQ ID NOS: 646
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 597
;   LENGTH: 25950
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-764-870-597

  Query Match             3.6%;  Score 35.6;  DB 10;  Length 25950;
  Best Local Similarity   51.2%;  Pred. No. 21;
  Matches   83;  Conservative    0;  Mismatches   79;  Indels    0;  Gaps    0;

Qy     118 AACACGCATCTTATAGAAACGTCCTATGATAGGTTGAAATCAAGAGAAATCACATTTCAG 177
           ||   | | ||  ||| |||     | |  |   | | |  || || |||| || |  ||
Db   18747 AAACTGGACCTAGTAGTAACAGTAAAAGTAATAGTAATAATAACAGCAATCTCAGTAAAG 18806

Qy     178 CAATACAGGGAAAATCTTGCTAAAGCAGGAGTTTTCCGATGGGTTACAAATATCCATGAA 237
             || | |     | | |  |||||| |  |  |||||  |    |    | |  ||| ||
Db   18807 AAACAAAATTCCAGTTTCACTAAAGAATTATCTTTCCCTTTTTATGTGCAAAAACATAAA 18866

Qy     238 CATAAAAGATATTACTATACCTTTGATAATTCATTACTATTT 279
            |||||||   |   |  | |    |  | | |||||||| ||
Db   18867 AATAAAAGGATTCTGTTGATCAAGCAGTAATGATTACTAATT 18908


Search completed: January  4, 2003, 05:22:23
Job time : 129.5 secs
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